

STIC Search Report Biotech-Chem Library

STIC Database Tracking Number: 154530

TO: Brian Kwon

Location: REM-3A74/3C70

Art Unit: 1614

Friday, May 27, 2005

Case Serial Number: 09/763616

From: Barb O'Bryen

Location: Biotech-Chem Library

Remsen 1a69

Phone: 571-272-2518

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barbara.obryen@uspto.gov

Search Notes

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SEARCH REQUEST FORM

Scientific and Technical Information Center

requester's Full Name: Brian In Unit: 1614 Phone Note all Box and Bldg/Room Location Phone State of the Son	Results	Format Preferred (circle): (PAPER)	DISK E-MAIL
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Database

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Length 37; 0; Indels

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Similarity 100. 37; Conservative

Query Match Best Local 8

Matches

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The present invention describes an agent used in the treatment or prophylaxis of a disease caused or exacerbated by MPV (mammalian papillomavirus) comprising a compound capable of reducing, inhibiting or otherwise decreasing the activity of a protein encoded by an MPV gene by facilitating disruption of a chelated metal cation domain present in the protein. An agent of the present invention can be used to treat cervical cancer or its HPV associated precursor lesions or other HPV associated cancers and/or warts. The present sequence represents a specifically claimed conserved motif sequence for a chelated zinc domain
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Polysulfide and dithionodisulfide agents, useful for the treatment or
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Human; secreted protein; diagnosis; cytostatic; immunosuppressive; antiHIV; antinflammatory; nootropic; neuroprotective; antiallergic; osteopathic; antiarthritic; antibacterial; antidiabetic; antiasthma; antipsoriatic; cardiant; gene therapy; cancer; neurological disorder; immune disease; inflammation; blood disorder; tumour.

Human secreted protein sequence encoded by gene 17 SEQ ID NO:287.

(first entry)

29-JUN-2000

AAY91614 standard; protein; 37

AAY91614

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human secreted proteins given in AAA26346 to AAA26458 encode the human secreted proteins given in AAY91451 to AAY91691. The human secreted proteins given in AAY91451 to AAY91691. The human secreted proteins given in AAY91451 to AAY91691. The human secreted proteins are based on the tissues and cells they are expressed in. Examples of the activities are: cytostatic; antidiabetic; immunosuppressive; antidiation; antidiated proteins are useful for preventing, treating or antidiated antidiated proteins are useful for preventing, treating or corresponding secreted proteins are useful for preventing, treating or pathological conditions, e.g. by protein or gene therapy. Also pathological conditions can be diagnosed by determining the amount of the proteins in a sample or by determining the presence of mutations in the polymucleotides, based on which tissues they are most highly expressed in, and include developing products for the diagnosis or treatment of cancer, tumoures, neurodegenerative disconders, developmental abnormalities and foetal deficiencies, blood discorders, discases of the immune system, autoimmune diseases, hepatic and renal diseases inflammation, allergies, Alzheimer's and behavioural disorders, cantioner, archimet, sinfections, AIDS, spinal cord injuries, transplant rejection, diabetes, asthma, sepsis, acne, psoriasis, cardiovascular disorders,
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Sequence 37 AA;

New isolated human genes and the secreted polypeptides they encode, useful for diagnosis and treatment of e.g. cancers, neurological disorders, immune diseases, inflammation or blood disorders.

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Komatsoulis GA, Rosen CA, Ruben SM, Duan R, Moore PA, Lafleur D, Wei Y, Ni J, Florence KA, Young PE, Brewer Soppet DR, Endress GA, Ebner R, Olsen HS, Mucenski M;

WPI; 2000-195282/17.

98US-0095486P. 98US-0095454P. 98US-0095455P.

05-AUG-1998;

06-AUG-1998 12-AUG-1998

98US-0096319P

(HUMA-) HUMAN GENOME SCI INC

99WO-US017130 98US-0094657P.

29-JUL-1999;

10-FEB-2000

WO200006698-A1. Homo sapiens.

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Matches
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reproductive disorders, gastrointestinal disorders, respiratory disorders and metabolic disorders. The proteins or polymucleotides can also be used as food additives or preservatives. The proteins are also useful for identifying their binding partners. AAA26337 to AAA26345 and AAY91450 are sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                              leukaemia; autoimmune disorder; allergy; cancer; cardiovascular disorder; respiratory disorder; ulmonary disorder; connective tissue disorder; skin disorder; CNS disorder; congented disorder; infectious disorder; gastrointestinal disorder; human; secreted protein.
                                                                                                                                                                                                                                                                                          anticonvulsant, antilipaemic; CNS; gynaecological; antiarthritic; antiathmatic; anti-HIV; virucide; endocrine; cytostatic; immunosuppressive; antiallergic; cardiovascular; respiratory; dermatological; antimicrobial; gastrointestinal; gene therapy; neurodegenerative disease; behavioral disorder; inflammatory condition;
                                                                                                                                                                                                                                                                                                                                          hyperproliferative disorder; Alzheimer's disease; Parkinson's disease;
Huntington's disease; metabolic disorder; Tay-Sach's disease;
Leash-Whan syndrome; reproductive disorder; immunological disorder;
arthritis; asthma; AlDS; andocrine disorder; immune disorder;
Hodgkin's lymphoma; haematopoietic disorder; muscular disorder;
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                                                                                                                          Novel human secreted protein fragment seqid 294.
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33; Mismatches 0;
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98US-0095454P.
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Wei Y;
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LAFLEUR D W.
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Best Local Similarity
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RUBEN S M.
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                                                               Sequence 37 AA;
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The invention describes an isolated human nucleic acid molecule (I)

Comprising a polynucleotide having a nucleotide sequence at least 954
identical to: a sequence polynucleotide fragment of ESQ ID NO: X or of
the cDNA sequence included in ATCC Deposit No: Z, which is hybridisable

CC the cDNA sequence included in ATCC Deposit No: Z, which is hybridisable to SEQ ID NO: X, not a sequence encoded by the cDNA

Sequence included in ATCC Deposit No: Z, which is hybridisable to SEQ ID

NO: X, having a biological activity. The nucleic acids and polypeptides,
pharmaceutical formulations and kits are useful in diagnosing and

treating neurodegenerative diseases states, behavioral disorders,
inflammatory conditions, hyperproliferative disorders (e.g. Alzheimer's

disease, Parkinson's diseases or Huntington's diseases), metabolic

disorders (e.g. Tay-Sach's disease or Leash-Nythan syndrome), reproductive

disorders (e.g. Tay-Sach's disease or Leash-Nythan syndrome), haematopoietic

cdisorders (e.g. Tay-Sach's disease or Leash-Nythan syndrome), haematopoietic

cdisorders and immune disorders (e.g. Hodgkin's lymphoma), haematopoietic

conditions afflicting connective tissue, skin disorders, allergy,
cancer, cardiovascular, respiratory or pulmonary disorders, allergy,
cancer, cardiovascular, respiratory or pulmonary disorders, alsorders,
conditions afflicting connective tissue, skin disorders, alsorders,
conditions afflicting connective tissue, skin disorders in fectious disorders and gastrointestinal

conditions afflicting connective tissue, skin disorders or conditions fragment of the invention. Note: This sequence does not appear in

the printed specification but is available in electronic format from the

cut figure at ftp.seqdata.uspto.gov/seqdata.html?
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New human secreted nucleic acid, useful for diagnosing and treating neurodegenerative, inflammatory, hyperproliferative, metabolic, reproductive, cardiovascular, respiratory or immunological disorders or
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10.8%; Pred. No. 6.8e+02;
ive 33; Mismatches 0;
                                                                                                                                                                                           Disclosure; SEQ ID NO 294; 234pp; English.
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99US-0128234P
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Best Local Similarity
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25-MAR-1999;
29-MAR-1999;
01-APR-1999;
06-APR-1999;
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05-MAR-1999;
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9US - 0128714P 9US - 0129845P 9US - 0120845P 9US - 0130449P 9US - 0130691P 9US - 01312448P 9US - 0132444P 9US - 0132444P 9US - 0132486P 9US - 0132481P 9US - 0132481P 9US - 0132481P 9US - 0132481P 9US - 0132481P 9US - 0132481P 9US - 0134211P 9US - 0134211P 9US - 0134211P 9US - 0134211P 9US - 0134211P 9US - 0134211P	990S-0136021P- 990S-0136021P- 990S-0137222P- 990S-0137222P- 990S-0137522P- 990S-0137522P- 990S-013864P- 990S-013864P- 990S-0139452P- 990S-0139452P- 990S-0139452P- 990S-0139458P- 990S-0139458P- 990S-0139456P- 990S-0139456P- 990S-0139456P- 990S-0139460P- 990S-0139460P- 990S-0139460P- 990S-0139460P- 990S-0139460P- 990S-0139460P- 990S-0139460P- 990S-0139460P-	905-0139817E 905-0139817E 905-01403318 905-01403184E 905-0140921E 905-0140921E 905-0141287E 905-0142154E 905-0142154E 905-014287E 905-01428019 905-01428019 905-0144086E 905-01443854E
3- APR - 1999 5- APR 1999 9- APR 1999	25. MAY - 1999; 27. MAY - 1999; 21. JUN - 1999; 03. JUN - 1999; 04. JUN - 1999; 06. JUN - 1999; 10. JUN - 1999; 14. JUN - 1999; 16. JUN - 1999; 17. JUN - 1999; 18. JUN - 1999;	1. Jun 1999 2. Jun 1999 3. Jun 1999 3. Jun 1999 4. Jun 1999 9. Jun 1999
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19-UU-199 19-UU-199 20-UU-199 20-UU-199 20-UU-199 21-UU-199 21-UU-199 21-UU-199 22-UU-199 22-UU-199 22-UU-199 23-UU-199 23-UU-199	23-JUL-19 22-JUL-19 22-JUL-19 27-JUL-19 27-JUL	, , , , , , , , , , , , , , , , , , ,

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The invention relates to a nucleic acid probe for measuring human gene expression, comprising any of the 27,400 fully defined nucleotide according at least 8 amino acids of any of the 6888 amino acid sequences in the specification, or their complements or fragments, and candidates under high stringency conditions to a nucleic acid molecule chybridises under high stringency conditions to a nucleic acid molecule acreased in human cells or tissues. Also included are a spatially.

Compression (comprising a plurality of single exon nucleic acid defressable set of single exon nucleic acid probes for measuring human cells or mucleic acid for the plurality of probes for steed above, where each of the plurality of probes for sequences or contaguous amino acids of any of the above mentioned amino acid contiguous amino acids of any of the above mentioned amino acid so contiguous amino acids of any of the above mentioned amino acid so contiguous amino acids of any of the above mentioned amino acid so contiguous amino acids of any of the above mentioned amino acid so contiguous amino acids of any of the above mentioned amino acid so contiguous amino acids of any of the above mentioned amino acid so contiguous amino acids of any of the above mentioned amino acid so contiguous amino acids of any of the above mentioned amino acid so contiguous amino acids of any of the above mentioned amino acid so contiguous amino acids and/or licensing single exon probes or microarrays to a customer destring to measure gene expression, a method of providing exonation acids and/or licensing single exon probes or microarrays. The probes may be used as tools for surveying crorage medium which contains a database having a plurality of records crorage medium which contains a database having a plurality of excords crorage medium which contains a database having a single exon probes or a used in identifying and characterising according to a probes are used in identifying and characterising alterrations in the genomic locatering spicious probes or a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peptide #273 encoded by probe for measuring cervical gene expression.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 69; DB 8; Length 47; 10.8%; Pred. No. 9.4e+02; ive 33; Mismatches 0; Indels
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26-MAY-2000; 2000US-0207456P.
30-JUN-2000; 2000US-00608408.
03-AUG-2000; 2000US-00632366.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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99US-0159295P.
99US-0159329P.
99US-01593310P.
99US-01596317P.
99US-0159638P.
99US-0160741P.
99US-016077P.
99US-016077P.
99US-016078P.
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99US-0161406P.
99US-0161359P.
99US-0161360P.
99US-0161361P.
99US-0161920P.
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Matches 4; Conserv
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surveying tissues.
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14-0CT-1
18-0CT-1
21-0CT-1
21-0CT-1
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21-0CT-1
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ABO59388
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AAM26246;
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                                                                                                                                                  The present invention relates to human single exon nucleic acid probes (SENP: see AAI10068-AAI28459). The present sequence is a peptide encoded by one such probe. The SENPs are derived from human Hela cells. The SENPs can be used to produce a single exon microarray, which can be used for measuring human gene expression in a sample derived from human cervical epithelial cells. By measuring gene expression, the probes are therefore useful in grading and/or staging of diseases of the cervix, notably cervical cancer. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human genome-derived single exon nucleic acid probes useful for analyzing gene expression in human fetal liver.
                                                                                                  Human genome-derived single exon nucleic acid probes useful for analyzing gene expression in human cervical epithelial cells.
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CIRCGGSVHVQSHSRLLTHSRNSIRIGRLIRSECLEC 47
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100.0%; Score 69; DB

Best Local Similarity 10.8%; Pred. No. 1e+0

Matches 4; Conservative 33; Mismatches
                                                                                                                                Claim 27; SEQ ID NO 18665; 487pp; English.
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                                       (MOLE-) MOLECULAR DYNAMICS INC.
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2000US-00608408.
2000US-00632366.
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2000US-0236359P.
2000GB-00024263.
21-SEP-2000; 2000US-0234687P.
         27-SEP-2000; 2000US-0236359P 04-OCT-2000; 2000GB-00024263
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                                                            Hanzel
                                                                                                                                                                                                                                                                    50 AA;
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30-JUN-2000;
03-AUG-2000;
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27-SEP-2000;
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                                                    The invention relates to a single exon nucleic acid probe for measuring human gene expression in a sample derived from human foctal liver. The single exon nucleic acid probes may be used for predicting, measuring and displaying gene expression in samples derived from human fetal liver. The present sequence is a poptide encoded by a single exon nucleic acid probe of the invention. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peptide #283 encoded by probe for measuring placental gene expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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Claim 27; SEQ ID NO 25419; 639pp + Sequence Listing; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       %; Score 69; DB 4;
; Pred. No. 1e+03;
33; Mismatches
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30-UUN-2000; 2000US-00608408.
03-AUG-2000; 2000US-00632566.
21-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0234587P.
04-OCT-2000; 2000GB-00024263.
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10.8%; P
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                                                                                                                                                                                                                                                                                                                                                                                                                                               4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               genetic disorder
                                                                                                                                                                                                                                                                                                                             Sequence 50 AA;
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from human breast and BT 474 cells. The method involves contacting the probes for measuring gene expression in a sample derived from human breast and BT 474 cells. The method involves contacting the probes with a collection of detectably labelled nucleic caids derived corner when the measuring the label bound to each probe of the microarray. The probes are useful for verifying the expression of regions of genomic DNA predicted to encode proteins. They are useful for gene discovery, and for determining predisposition and/or prognosing breast disease. Gene expression analysis is useful for assessing the toxicity of chemical agents on cells. The microarray of this invention presents a far greater diversity of probes for measuring gene expression, with far less has than expressed sequence tag microarrays. The method is suitable for rapid production of functional information from genomic sequence. The present sequence is a peptide cencoded by a single exon nucleic acid probe of the invention. Note: The sequence data for this patent did not form part of the printed sequence but was obtained in electronic format directly from WIPO
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                                                                                                                                                                                                                                                                                                               Human peptide #265 encoded by breast cell single exon nucleic acid probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New spatially-addressable set of single exon nucleic acid probes, useful for measuring gene expression in sample derived from human breast, comprises number of single exon nucleic acid probes.
                                                                                                                                                                                                                                                                                                                                                    Human; microarray; single exon probe; gene expression; breast; disease;
                                    Gaps
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                                Indels
                                                                                            11 CIRCCGSVHVQSHSRLLTHSRNSIRIGRLIRSECLEC 47
                                  0
                                                                     1e+03;
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               Pred. No. 1e+C
; Mismatches
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10.8%; Pic.
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21-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0236359P.
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2000US-0207456P.
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                                                                                                                                                                                                                                                                            (first entry)
                                  Conservative
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           Best Local Similarity
Matches 4; Conserv
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                                                                                                                                                                                                                                                                                                                                                                           cancer.
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The present invention relates to single exon nucleic acid probes for measuring human gene expression in a sample derived from human heart (see ABA21535-ABA41305). The present sequence is a protein encoded by one such probe. The probes may be used for predicting, measuring and displaying gene expression in samples derived from the human heart via microarrays. By measuring gene expression, the probes are useful for predicting, diagnosing, grading, staging, monitoring and prognosing diseases of the human heart and vascular system e.g. cardiovascular disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 hypertension, cardiac arrhythmias and congenital heart disease. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                       Protein #266 encoded by probe for measuring heart cell gene expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Single exon nucleic acid probes for analyzing gene expression in human
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                                                                                                                                                                                                                                     Human, gene expression, heart, microarray, vascular system, cardiovascular disease, hypertension, cardiac arrhythmia, congenital heart disease.
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37
                  11 CIRCCGSVHVQSHSRLLTHSRNSIRIGRLIRSECLEC 47
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100.0%; Score 69; DB 4, 10.8%; Pred. No. 1e+03; iive 33; Mismatches
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                                                                                                           ABB18267 standard; protein; 50
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27-SEP-2000; 2000US-0236359P.
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30-JUN-2000; 2000US-00608408
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Best Local Similarity
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                                                                             RESULT 10
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RESULT 11 AAM65971

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probes which are derived from genomic sequences expressed in the human brain. They can be used to measure gene expression in brain cell samples, which may enable the diagnosis and improved treatment of nervous system diseases such as Alzheimer's disease, multiple sclerosis, schizophrenia, epilepsy and cancers. The present sequence is a protein encoded by one of the probes of the invention
                                                                                                                                                                                                                                                                                                          Single exon nucleic acid probes for analyzing gene expression in human
                                                                                                                                                                                                                                                                                                                                                                                              The present invention provides a number of single exon nucleic acid
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hypercholesterolaemia, coronary heart disease.
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2000US-00608408
2000US-00632366
2000US-0234687P
2000US-0234687P
2000US-023458P
2000US-0236359P
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                                                                                                                                    21-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0236359P.
04-OCT-2000; 2000GB-00024263.
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                                                                2000US-0180312P.
2000US-0207456P.
2000US-00608408.
                                30-JAN-2001; 2001WO-US000667.
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21-SEP-2000; 2000US-0234687P.
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                                                                                                                                                                                                                                      Penn SG, Hanzel DK,
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 50 AA;
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03-AUG-2000; 2
21-SEP-2000; 2
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                                                                                                    30-JUN-2000;
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                                                                04-FEB-2000;
26-MAY-2000;
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09-AUG-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     probes which are derived from genomic sequences expressed in the human bone marrow. They can be used to measure gene expression in bone marrow samples, which may enable the improved diagnosis and treatment of cancers such as lymphoma, leukaemia and myeloma. The present sequence is a protein encoded by one of the probes of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; brain expressed exon; gene expression analysis; probe; microarray; Alzheimer's disease; multiple sclerosis; schizophrenia; epilepsy; cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 genome-derived single exon nucleic acid probes useful for analyzing expression in human bone marrow.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       present invention provides a number of single exon nucleic acid
                                                                                                                                  Human, bone marrow expressed exon, gene expression analysis; probe, microarray; cancer; leukaemia; lymphoma; myeloma.
                                                                                                    Human bone marrow expressed probe encoded protein SEQ ID NO: 26277
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AAM65971 standard; protein; 50 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MOLE-) MOLECULAR DYNAMICS INC
                                                                                                                                                                                                                                                                                                                                              2000US-0207456P.
2000US-00608408.
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2000US-0234687P,
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                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-488900/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 50 AA;
                                                                                                                                                                                                                          WO200157276-A2.
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03-AUG-2000;
21-SEP-2000;
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                                                                   06-NOV-2001
                                                                                                                                                                                                                                                            09-AUG-2001
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                                 AAM65971;
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Human gene

Matches

RESULT 12 AAM5359.

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Gaps

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Indels

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Claim 27; SEQ ID NO 10323; 322pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-114183/15.
                                                                                                                                                                                                                            Sequence 50 AA;
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26-MAY-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                        ABG35618;
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                                                                                                                    The invention relates to a single exon nucleic acid probe (SENP) (I) for measuring human gene expression in a sample derived from human adult liver, comprising one of 13109 defined nucleotide sequences given in the specification (or complements/ fragments). The probe hybridises at high stringency to a nucleic acid modecule expressed in the human adult liver. (I) may be used for predicting, measuring and displaying gene expression in samples derived from human adult liver. The genes identified may be involved in genetic liver diseases such as cirrhosis, hyperlipoproteinaemia, hyperlipidaemia and hypercholesterolaemia which is associated with coronary heart disease. ABG47348-ABG59930 represent human liver single exon encoded peptides of the invention. Note: The sequence information for this patent doses not appear in the printed specification but was obtained in electronic format directly from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                      Human genome-derived single exon nucleic acid probes useful for analyzing gene expression in human adult liver.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel single exon nucleic acid probe used to measuring gene expression in a human breast.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peptide #265 encoded by probe for measuring human breast gene expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Probe; human; breast disease; breast cancer; development disorder; inflammatory disease; proliferative breast disease; non-carcinoma tumour.
                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                  4; Length 50;
                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
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CIRCCGSVHVQSHSRLLTHSRNSIRIGRLIRSECLEC 47
                                                                                                                                                                                                                                                                                                                                                                                                                      %; Score 69; DB 4;
; Pred. No. 1e+03;
33; Mismatches
                                                                                             Claim 27; SEQ ID NO 26284; 658pp; English
                                                                                                                                                                                                                                                                                                         ftp.wipo.int/pub/published_pct_sequences
 Rank DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rank DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAM01583 standard; protein; 50 AA
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2000US-00608408.
2000US-00632366.
2000US-0234687P.
2000US-0236359P.
 Chen W,
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 Hanzel DK,
                           WPI; 2001-488898/53
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Best Local Similarity
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03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
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The present invention relates to novel single exon nucleic acid probes (see AAI00010-AAI10067). The present sequence is a peptide encoded by one auch probe. The probes are useful for measuring human gene expression in a human breast sample, where the probe hybridises at high stringency to a predicting, diagnosing, grading, staging, monitoring and prognosing diseases of the human breast, particularly those diseases with polygenic actiology. The diseases include: breast cancer, disorders of development, inflammatory diseases of the breast, fibrocystic changes, proliferative breast disease and non-carcinoma tumours. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human peptide encoded by genome-derived single exon probe SEQ ID 25283.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hermansky-Pudlak syndrome; sarcoidosis; pulmonary haemosiderosis; pulmonary histiocytosis; lymphangioleiomyomcosis; Karagener syndrome; pulmonary alveolar proteinosis; fibrocystic pulmonary dysplasia; primary ciliary dyskinesis; pulmonary hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Spatially-addressable set of single exon nucleic acid probes, used to
                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; single exon probe; asthma; lung cancer; COPD; ILD; chronic Obstructive pulmonary disease; intersetitial lung disease; familial idiopathic pulmonary fibrosis; neurofibromatosis; tuberous sclerosis; Gaucher's disease; Niemann-Pick disease;
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                                                                                                                                                                                                                                                                                                                        DB 4; Length 50;
                                                                                                                                                                                                                                                                                                                                                             0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                            measure gene expression in human lung samples.
                                                                                                                                                                                                                                                                                                                                         1e+03;
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; Pred. No. 1e+0
33; Mismatches
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                                                                                                                                                                                                                                           ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABG35618 standard; peptide; 50
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2000US-0207456P.
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03-AUG-2000; 2000US-00632366
21-SEP-2000; 2000US-0234687P
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Best Local Similarity 10.8*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hyaline membrane disease.
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The invention relates to a spatially-addressable set of single exon nucleic acid probes for measuring gene expression in a sample derived from thuman lung comprising single exon nucleic acid probes having one of 12614 nucleic acid sequences mentioned in the specification, or their complements or the 123187 open reading frames derived from the 12614 probes. Also included are a matcroartay comprising the novel set of probes which phyridises at high stringency to a nucleic acid expressed in the human lung; measuring gene expression in a sample derived from human lung; comprising (a) contacting the array with a sample derived from human lung man, and included are a matcroartay comprising (b) measuring the labeled nucleic acids derived from human lung man, a eukaryotic genome, comprising (a) detecting specific hybridiatation of detectably and (b) measuring the labeled nucleic acids derived from human lung man, a comprising (a) detecting specific hybridiatation of detectably predicting at least one exon from genomic sequences of the eukaryote; and (b) detecting specific hybridiatation of detectably labeled nucleic acids from entationed microarray, assigning exons to a single exon probe having a fragment identical to the predicted exon, the probe is included the above mentioned microarray, assigning exons to a single exon microarray having the exons from genomic sequence by the method microarray having the exons in the tissues and/or cell types using hybridiation to a single exon microarrays having a probe with the exon where a common pattern of the exons should be assigned to a single gene, a peptide comprising on the exons in the tissues and/or cell types using hybridiation to a single exon in grobes/open reading frames (ORFR). The probes are used for gene expression of the exons in a denomery diseases such as astimate the probes where a common pattern of alsease into himmany diseases such as astimate to the exons common pattern of derived mRNN and for the exons in a gene hypothesis, pulmonary diseases (DDP), inters

Gaps ö Query Match
100.0%; Score 69; DB 5; Length 50;
Best Local Similarity 10.8%; Pred. No. 1e+03;
Matches 4; Conservative 33; Mismatches 0; Indels

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11 CIRCCGSVHVQSHSRLLTHSRNSIRIGRLIRSECLEC 47

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Search completed: May 27, 2005, 00:32:23 : 165 secs Job time

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GENERAL INFORMATION:
APPLICANT: Goodearl, Andrew; Stroobant, Paul;
APPLICANT: Goodearl, Andrew; Stroobant, Michael; Marchioni, Mark;
APPLICANT: Chen, Maio Su; Hiles, Ian
APPLICANT: Chen, Maio Su; Hiles, Ian
TITLE OF INVENTION: Glial Mitogenic Factors, Their
TITLE OF INVENTION: Preparation and Use
NUMBER OF SEQUENCES: 184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-036-555B-40
; Sequence 40, Application US/08036555B
; Patent No. 5530109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                             US-09-489-847-294
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LENGTH: 37
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Sequence 40, Appl
Sequence 19, Appl
Sequence 38, Appl
Sequence 39, Appl
Sequence 91, Appl
Sequence 8, Appl
Sequence 8, Appl
Sequence 34381, A
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Sequence 302, App
Sequence 183, App
Sequence 183, App
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                                                              May 27, 2005, 00:27:20 ; Search time 42 Seconds (without alignments) 65.762 Million cell updates/sec
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Patent No. 5258
Patent No. 5258
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/cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
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/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
                                                                                                                                   Compugen Ltd.
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US-08-036-555B-40
US-08-269-569-40
US-08-469-526A-40
US-08-745-501A-40
US-08-735-021-40
US-08-735-021-40
US-08-735-021-40
US-08-735-06846A-40
PCT-US94-05083C-40
PCT-US94-05083C-40
US-08-230-46
US-08-691-814B-38
US-08-691-814B-39
US-08-691-814B-30
US-08-691-814B-30
US-08-691-814B-30
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US-09-402-181B-302
US-09-721-456-302
5258287-15
5258287-15
GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compug
                                                                                                                                                                                                              Total number of hits satisfying chosen parameters:
                                                                                                                                                                                          513545 segs, 74649064 residues
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                                                                                                                                                                                                                                                                    Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                            OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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                                                                                                                                                                                                                                     Minimum DB seq length: 0
Maximum DB seq length: 200000000
                                                                                                            US-09-763-616-2
69
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90, Appl
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13, Appl
33917, A
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US-08-476-509B-46
US-10-125-062-4
US-09-252-991A-23832
US-08-852-666-14
US-09-471-276-941
US-09-471-276-941
US-09-471-276-941
US-09-270-776-121
US-09-854-864-13
US-09-270-776-742140
US-09-854-864-13
US-09-270-776-743917
US-09-270-777-4218
US-09-860-793-5
US-09-419-381-90
US-09-419-381-90
US-08-406-248-6
US-08-406-248-6
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US-08-406-248-6
US-08-606-248-6
US-08-606-248-6
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TITLE OF INVENTION: 98 Human Secreted Proteins
FILE REFREENCE: PSO31P1
CURRENT PILING DATE: 2000-01-24
EARLIER APPLICATION NUMBER: US/09/489,847
CURRENT FILING DATE: 1999-07-29
EARLIER PILING DATE: 1999-07-29
EARLIER PILING DATE: 1999-07-30
EARLIER FILING DATE: 1998-07-30
EARLIER FILING DATE: 1998-07-30
EARLIER FILING DATE: 1998-08-05
EARLIER FILING DATE: 1998-08-05
EARLIER FILING DATE: 1998-08-05
EARLIER FILING DATE: 1998-08-15
EARLIER PILING DATE: 1998-08-06
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-09-382-616A-1
                                                                                                                                                                                                                                                                                                                ALIGNMENTS
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MEDIUM TYPE: Diskette, 5.25 inch, 360 kb storage COMPUTER: IBM OPERATING SYSTEM: PC-DOS
                                                                                          SOFTWARE: Wordperfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/469,569
FILING DATE: 06-JUN-1995
CLASSIFCATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: TSAI, Christine H.
REGISTRATION NUMBER: 34,266
REFERENCE/DOCKET NUMBER: LUD
TELECOMMUNICATION:
TELECOMMUNICATION:
(212) 688-9200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Felfe & Lynch
STREET: 805 Third Avenue
CITY: New York City
STATE: New York
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: 1BM
OPERATING SYSTEM: PC-DOS
SOFTWARE: Wordperfect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (212) 838-3884
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-249-322A-40
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US-08-469-569-40
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Sequence 40, Application US/08469569

Sequence 40, Application US/08469569

GENERAL INPORMATION:

APPLICANT: Goodearl, Andrew; Stroobant, Paul;

APPLICANT: Minghetti, Luisa; Waterfield, Michael; Marchioni, Mark;

APPLICANT: Chen, Maio Su; Hiles, Ian

TITLE OF INVENTION: Glal Mitogenic Factors, Their

TITLE OF INVENTION: Preparation and Use

NUMBER OF SEQUENCES: 184

CORRESPONDENCE ADDRESS:

ADDRESSE: Pelfe Lynch

STREET: 805 Third Avenue

CITY: New York City

STREET: New York

STREET: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 360 kb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 CGCCKCCRTTCACRCAGAAGGTCTTCTCCTTCTCAGC 39
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                                                                                                                                                                                                                                                                                          CUKKENI PEPLIANIN DAIA:
CUKKENI PEPLIANIN DAIA:
FILING DATE: 24-MAR-193
FILING DATE: 24-MAR-193
FRIOR APPLICATION DAIA:
APPLICATION NUMBER: 07/965,173
FILING DATE: 23-OCT-1992
FRIOR APPLICATION DAIA:
APPLICATION NUMBER: 07/940,389
FILING DATE: 03-SEP-1992
PRIOR APPLICATION DAIA:
APPLICATION NUMBER: 07/907,138
FILING DATE: 30-UN-1992
FILING DATE: 03-APRIL-1992
PRIOR APPLICATION DAIA:
APPLICATION NUMBER: 07/863,703
FILING DATE: 03-APRIL-1992
PRIOR APPLICATION DAIA:
APPLICATION NUMBER: 07/863,703
FILING DATE: 10-APRIL-1992
FILING DATE: 10-APRIL-1992
ATONNEY AGENT THORNAMINER: 0.K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Tsai, Christine H.
REGISTATION NUMBER: 34,266
REFERENCE, DOCKET NUMBER: LUD 5250.4
FELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (212) 688-9200
TELEFAX: (212) 838-3884
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                       ADDRESSEE: Felfe & Lynch
STREET: 805 Third Avenue
CITY: New York City
                                                                                                                                                                                                           COMPUTER: IBM
OPERATING SYSTEM: PC-DOS
                                                                                                                                                                                                                                                      SOFTWARE: Wordperfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/
       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TOPOLOGY: linear
US-08-036-555B-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              amino acid
                                                                                              New York
                                                                                                                    USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
                                                                                            STATE: New COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 10022
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Gaps
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Patent No. 5716930
GENERAL INFORMATION:
APPLICANT: Goodearl, Andrew, Stroobant, Paul;
APPLICANT: Minghetti, Luisa, Waterfield, Michael; Marchioni, Mark;
APPLICANT: Chen, Maio Su; Hiles, Ian
IITLE OF INVENTION: Glial Mitogenic Factors, Their
TITLE OF INVENTION: Preparation and Use
NUMBER OF SEQUENCES: 184
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIF: 10022
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 360 kb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CIALING DATE: 00-1001-1995
CIASSIFICATION 15-30
PRIOR APPLICATION DATA: 08 4036,555
FILING DATE: 24-MAR-1993
APPLICATION NUMBER: 07/965,173
FILING DATE: 23-0CT-1992
PRIOR APPLICATION NUMBER: 07/940,389
FILING APPLICATION DATE: 07/940,389
FILING DATE: 30-10N-1992
PRIOR APPLICATION NUMBER: 07/907,138
FILING DATE: 30-JUN-1992
PRIOR APPLICATION NUMBER: 07/863,703
FILING DATE: 03-APPLIL-1992
PRIOR APPLICATION NUMBER: 07/863,703
FILING DATE: 03-APRLIL-1992
PRIOR APPLICATION NUMBER: 07.863,703
FILING DATE: 10-APPLIL-1991
ATTORNEY/AGENT INFORMATION: 07/907
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Query Match 100.0%; Score 69; DB 1; Length 39; Best Local Similarity 10.8%; Pred. No. 1.18+02; Matches 4; Conservative 33; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Goodearl, Andrew
APPLICANT: Goodearl, Paul
APPLICANT: Minghetti, Luisa
APPLICANT: Minghetti, Luisa
APPLICANT: Minghetti, Luisa
APPLICANT: Marchionni, Mark
APPLICANT: Marchionni, Mark
APPLICANT: Chen, Mario
ITTLE OF INVENTION: GLIAL MITOGENIC FACTORS, THEIR
ITTLE OF INVENTION: GLIAL MITOGENIC FACTORS, CORRESPONDENCE ADDRESS:
ADDRESSEE: Clark & Elbing LLP
STREET: 176 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3 GCCKCCRITCACRCAGAAGGTCTICTCCTTCTCAGC 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PULGATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 08/036,555
FILING DATE: 24-MAR-1993
FILING DATE: 23-OCT-1992
APPLICATION NUMBER: 07/96,173
FILING DATE: 03-OCT-1992
APPLICATION NUMBER: 07/90,138
FILING DATE: 03-UN-1992
APPLICATION NUMBER: 07/90,7138
FILING DATE: 03-APRIL-1992
APPLICATION NUMBER: 07/86,703
FILING DATE: 10-APR-1991
ATTORNEY AGENT INFORMATION:
NAME: Bisker-Erady, Kristina
REGISTRATION NUMBER: 39,109
TELEROMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM COMPUTER: OPERATING SYSTEM: Windows95
OPERATING SYSTEM: Windows95
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/734,591A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 02110
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
COMPUTER: IBM Compatible Pentium
COMPUTER: Mindows95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 40, Application US/08734591A Patent No. 5854220
                  APPLICATION NUMBER: US/08, FILING DATE: 06 June 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 39
    CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-469-526A-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Boston
Massachusetts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: U.S.A. ZIP: 02110
                                            FILING DATE: 06
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-734-591A-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: E
STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GLIAL MITOGENIC FACTORS, THEIR PREPARATION AND USE 187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3 CGCCKCCRTTCACRCAGAAGGTCTTCTCTCTCTCTCAGC 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM Compatible OPERATING SYSTEM: DOS SOFTWARE: FastSEQ for Windows Version 2.0
                                                               PICRASIFICATION: 435
PRIOR APPLICATION NUMBER: 08/036,555
FILING DATE: 24-MAR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/965,173
FILING DATE: 23-OCT-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/940,389
FILING DATE: 03-SEP-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/90,138
FILING DATE: 30-JUN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/863,703
FILING DATE: 03-APRIL-1992
PRIOR APPLICATION NUMBER: 07/863,703
FILING DATE: 10-APRIL-1991
ATTORNEY/AGENT INFORMATION:
NAME: TBAL CATION NUMBER: 14.6
REFIERENCE/DOCKET NUMBER: 14.26
REFERENCE/DOCKET NUMBER: LUD 250.4
TELEPHANE: (212) 688-9200
TELEPHANE: (212) 688-9200
TELEPHANE: (212) 688-9200
TELEPHANE: 70-MARCHISTICS:
    .оли Е.R.: US/08/249,322A
26-MAY-1994
V: A.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: Clark & Elbing LLP STREET: 176 Federal Street CITY: Boston STATE: MA COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 40, Application US/08469526A Patent No. 5792849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Stroobart, Paul
APPLICANT: Minghetti, Luisa
APPLICANT: Marchionni, Mark
APPLICANT: Chen, Maio Su
APPLICANT: Chen, Maio Su
APPLICANT: Chen, Maio Su
TITLE OF INVENTION: CLIAL MITOGER
TITLE OF INVENTION: PREPARATION INVERED ON SEQUENCES: 187
CORRESONDENCE ADDRESS:
ADDRESSEE: Clark & Elbing LIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Goodearl, Andrew
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                       APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TOPOLOGY: linear
US-08-249-322A-40
                                          FILING DATE: 26 CLASSIFICATION:
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: HILES, IAN
APPLICANT: HILES, IAN
TITLE OF INVENTION: PREPARATION AND USE
FILE REPERENCE: 04585/00200L
CURRENT APPLICATION NUMBER: US/08/735,021B
CURRENT APPLICATION NUMBER: US/08/735,021B
CURRENT APPLICATION NUMBER: 08/412,065
EARLIER APPLICATION NUMBER: 08/412,065
EARLIER PILING DATE: 1995-06-06
EARLIER PILING DATE: 1995-06-05
EARLIER PILING DATE: 1993-03-24
EARLIER PILING DATE: 1992-06-33
EARLIER FILING DATE: 1992-06-33
EARLIER FILING DATE: 1992-06-33
EARLIER FILING DATE: 1992-06-33
EARLIER FILING DATE: 1992-06-30
EARLIER FILING DATE: 1992-06-30
EARLIER PELICATION NUMBER: 07/907,138
EARLIER FILING DATE: 1992-06-30
EARLIER PELICATION NUMBER: 07/907,138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3 CGCCKCCRITCACRCAGAAGGICITCTCCCTICTCAGG 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 69; DB 3;
10.8%; Pred. No. 1.1e+02;
tive 33; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1%; Score 69; DB 2;
5; Pred. No. 1.1e+02;
33; Mismatches 0
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/984,085
FILING DATE: 0.1-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/951,747
FILING DATE: 25-SEP-1992
FILING DATE: 25-SEP-1992
FILING DATE: 07/927,337
APPLICATION NUMBER: 07/927,337
FILING DATE: 10-AUG-1992
ATTORNEY/AGENT INFORMATION:
NAME: CLark, Paul T.
REGISTRATION NUMBER: 30.162
REFERENCE/DOCKET NUMBER: 04585/017004
TELEPHONE: (617) 542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 40, Application US/08735021B Patent No. 6194377 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
10.8%; F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: WATERFIELD, MICHAEL
APPLICANT: MARCHIONNI, MARK
APPLICANT: CHEN, MARIO S.
APPLICANT: CHEN, MARIO S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 10.8*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: GOODEARL, ANDREM
APPLICANT: STROOBANT, PAUL
APPLICANT: MINGHETTI, LUISA
                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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; ORGANISM: Bos taurus
US-08-735-021-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best_Local Similarity
Matches 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TOPOLOGY: linear
US-08-469-660-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           JS-08-735-021-40
                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 40
LENGTH: 39
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0
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100.0%; Score 69; DB 2; Length 39;
Best Local Similarity 10.8%; Pred. No. 1.1e+02;
Matches 4; Conservative 33; Mismatches 0; Indels
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Patent No. 5876973

GENERAL INFORMATION:
APPLICANT: GWYDNE, David I.; Marchionni, Mark;
APPLICANT: GWYDNE, David I.; Marchionni, Mark;
APPLICANT: GWBUTHON: INHIBITORS OF CELL PROLIFERATION, TITLE OF INVENTION: ITHEIR PREPARATION AND USE NUMBER OF SEGURDICES: 184

CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 225 Franklin Street
CITY: Boston
CITY: Bassachusetts
ZIP: 0211-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 360 kb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3 CGCCKCCRTTCACRCAGAAGGTCTTCTCCCTTCTCAGC 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 39,109
REFERENCE/DOCKET NUMBER: 04585/00200P
IELECOMMINICATION INFORMATION:
TELEPHONE: (617) 428-0200
                                                                                    PALLOK APPLICATION DATA:
PAPLICATION NUMBER: 08/036,555
PRIOR APPLICATION DATA: 1993
APPLICATION DATA: 1993
APPLICATION NUMBER: 07/965,173
FILING DATE: 23-0CT-1992
PRIOR APPLICATION DATA: 07/940,389
FILING DATE: 03-5EP-1992
PRIOR APPLICATION DATA: 07/907,138
FILING DATE: 30-JUN-1992
PRIOR APPLICATION DATA: 07/907,138
FILING DATE: 30-JUN-1992
PRIOR APPLICATION DATA: 07/863,703
FILING DATE: 03-APR-1992
PRIOR APPLICATION DATA: 07/863,703
FILING DATE: 10-APR-1992
PRIOR APPLICATION NUMBER: UK 91 07566.3
ATTORNEY/AGENT INFORMATION:
NAME: Bieker-Brady, Kristina
REGISTATION NUMBER: 39,109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: PC-DOS
SOGTWARE: Wordperfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/469,660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/011,396
FILING DATE: 29-JAN-1993
                     08/470,335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        E: Fish & Richardson
225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           40:
                                           FILING DATE: 06-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (617) 428-7045
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-734-591A-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-469-660-40
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APPLICANT: Goodearl, Andrew
APPLICANT: Stroobant, Paul
APPLICANT: Minghetti, Luisa
APPLICANT: Materfield, Michael
APPLICANT: Waterfield, Michael
APPLICANT: Waterfield, Michael
APPLICANT: Ghen, Marchionni, Mark
APPLICANT: Chen, Marchionni, Mark
TITLE OF INVENTION: GLIAL MITOGENIC FACTORS, THEIR
TITLE OF INVENTION: PREPARATION AND USE
NUMBER OF SEQUENCES: 187
CORRESPONDENCE ADDRESS:
ADDRESSEE: Clark F. T.
                         |::|::::::::|::|
CGCCKCCRTTCACRCAGAAGGTCTTCTCCCTTCTCAGC 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/734,664A
FILING DATE: 22-0CT-1996
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/249,322
FILING DATE: 26-MAY-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/036,555
FILING DATE: 24-MAR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/965,173
FILING DATE: 23-0CT-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/940,389
FILING DATE: 03-SEP-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/940,389
FILING DATE: 03-SEP-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/963,703
FILING DATE: 03-APR-1992
PRIOR APPLICATION NUMBER: UK 91 07566.3
FILING DATE: 03-APR-1991
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: US 91 07566.3
FILING DATE: 03-APR-1991
ATTORNEY/AGENT INFORMATION:
NAME: Bieker-Brady, Kristina
REGISTRATION NUMBER: 39,109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04585/00200J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 02110
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
COMPUTER: IBM Compatible Pentium
OPERATING SYSTEM: Windows95
SOFTWARE: FastSeq Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 04
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (617) 428-0200
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TELEFAX: (617) 428-7045
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Massachusetts: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TOPOLOGY:
US-08-734-664A-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
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Sequence 40, Application PC/TUS9506846A

Sequence 40, Application PC/TUS9506846A

PERREAL INFORMATION:

APPLICANT: Goodearl, Andrew David; Stroobant, Paul;

APPLICANT: Minghetti, Luiss; Waterfield, Michael; Marchioni, Mark;

APPLICANT: Chen, Maio Su; Hiles, Ian
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  Length 39;
                                                       0; Indels
                                                                                                                                             3 CGCCKCCRTTCACRCAGAAGGTCTTCTCCCTTCTCAGC 39
                                                                                                           3 CGCCKCCRITCACRCAGAAGGICITCICCTICTCTCAGG 39
Query Match

100.0%; Score 69; DB 3;

Best Local Similarity 10.8%; Pred. No. 1.1e+02;

Matches 4; Conservative 33; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 225 Franklin Street
CITY: Boston
STATE: Massachusetts
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 360
COMPUTER: IBM
                                                                                                                                                                                                                                                                                                                                                APPLICANT: Robert Sklar, Mark Marchionni,
APPLICANT: Bavid I. Gwynne
TITLE OF INVENTION: METHODS FOR ALTERING
TITLE OF INVENTION: MUSCLE CONDITION
NUMBER OF SEQUENCES: 185
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04585/028WO1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: PC-DOS
SOFTWARE: Wordperfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/05083C
FILING DATE: 06-MAY-94
CLASSIFICATION:
                                                                                                                                                                                                                                          RESULT 10
PCT-US94-05083C-40
; Sequence 40, Application PC/TUS9405083C
; GENERAL INFORMATION:
Darbort Sklar, Mark Marchi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/209,204
FILING DATE: 08-MAR-94
PRIOR APPLICATION NUMBER: 08/059,022
FILING DATE: 06-May-93
ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 04585/02
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3: Fish & Richardson
225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity 10.8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; TOPOLOGY: linear
PCT-US94-05083C-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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GENERAL INCRMATION:
APPLICANT: Bard, Jonathan A.
APPLICANT: Bard, Jonathan A.
TITLE OF INVENTION: DNA ENCODING GALANN GALR3 RECEPTORS AND
TITLE OF INVENTION: USES THEREOF
NUMBER OF SEQUENCES: 59
CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooper & Dunham LLp
STREET: 1185 Avenue of The Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 100.0%; Score 69; DB 3; Length 43; Best Local Similarity 10.8%; Pred. No. 1.38+02; Matches 4; Conservative 33; Mismatches 0; Indels
                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US 60/022,997
FILING DATE: US 60/022,997
FILING DATE: US 60/038,109
FILING DATE: US 60/038,109
FILING DATE: US-FEB-1997
ATTORNEY/AGENT INFORMATION:
NAME: Bak, MATY E.
REFERENCE/ODCKET NUMBER: 31,215
REFERENCE/ODCKET NUMBER: 31,215
REFERENCE/ODCKET NUMBER: 31,215
REFERENCE/ODCKET NUMBER: 31,215
TELEFPAX: 215-540-9200
NUMBER OF SEQUENCES: 47
CARRESPONDENCE ADDRESS:
ADDRESSER: Howson and Howson STREET: Spring House Corporate Cutr., PO Box 457
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4 CSRCTNILREPVCLGGCEHIFCSNCVSDCIGTGCPVC 40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/900,230
FILING DATE: 23-UUL-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-900-230-46
; Sequence 46, Application US/08900230
; Patent No. 6329197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 43 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
TOPOLOGY: unknown
MOLECULE TYPE: protein
                                                                                                   CITY: Spring House
STATE: Pennsylvania
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 23-JUL-
CLASSIFICATION: 435
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COUNTRY: U.S.A.
ZIP: 11036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-230-196-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19, Application US/09230196
Patent No. 6307035
GENERAL INFORMATION:
APPLICANT: Rauscher III, Frank J.
APPLICANT: Jensen, David E.
TILLE OF INVENTION: BRCA1 Associated Protein (BAP-1) and
TITLE OF INVENTION: Uses Therefor
  Glial Mitogenic Factors, Their
Preparation and Use
178
                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 360 kb storage
COMPUTER: IBM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3 CGCCKCCRTTCACRCAGAAGGTCTTCTCCTTCTCAGG 39
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APPLICATION NUMBER:
APPLICATION NUMBER:
FILING DATE: 24-MAR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 23-OCT-1992
PRIOR APPLICATION DATE:
APPLICATION NUMBER:
FILING DATE: 03-SEP-1992
PRIOR APPLICATION DATE:
APPLICATION NUMBER: 07/940,389
FILING DATE: 03-SEP-1992
PRIOR APPLICATION NUMBER: 07/907,138
FILING DATE: 30-JUN-1992
PRIOR APPLICATION NUMBER: 07/863,703
FILING DATE: 03-APRIL-1991
PRIOR APPLICATION NUMBER: 07.863,703
FILING DATE: 10-APRIL-1991
ATTORNEY/AGENT INFORMATION:
NAWE: HARSON NUMBER: 30,946
REPERRENCE/DOCKET NUMBER: 30,946
REPERRENCE/DOCKET NUMBER: 105-00000
                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM: PC-DOS
SOFTWARE: Wordperfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/06846A
FILING DATE: 25-MAY-1995
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
FILING DATE: 26-MAY-1994
CLASSIFICATION:
                                                                 CORRESPONDENCE ADDRESS:
ADDRESSE: Felfe & Lynch
STREET: 805 Third Avenue
CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (212) 838-3884
INFORMATION FOR SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (212) 688-9200
  TITLE OF INVENTION: 11
TITLE OF INVENTION: 11
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TOPOLOGY: linear
PCT-US95-06846A-40
                                                                                                                                                                                                          USA
                                                                                                                                                                                                       COUNTRY: US
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Search completed: May 27, 2005, 00:36:54 Job time: 43 secs
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APPLICANT: Tomasetto, Catherine
APPLICANT: Basset, Paul
APPLICANT: Byrne, Jennifer
TITLE OF INVENTION: Isolated Nucleic Acid Molecules Useful
TITLE OF INVENTION: as Leukemia Markers and in Breast Cancer Prognosis
NUMBER OF SEQUENCES: 124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;
0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Sterne, Kessler, Goldstein & Fox P.L.L.C. STREET: 1100 New York Ave, NW, Suite 600 CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY:

STATE: DC

COMPUTRY: USA

ZIP: 20005-3934

ZUP: 20005-3934

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: Ploppy disk

SOFTWARE: Patentin Release #1.0, Version #1.30

SOFTWARE: Patentin Release #1.0, Version #1.30

CURSTILICATION DATA:

APPLICATION NUMBER: US/08/691,814B

FILING DATE: 31-UUL-1996

CLASSIFICATION NUMBER: US 60/002,183

FILING DATE: 09-AUG-1995

ATTORNEY/AGENT INFORMATION:

NAME: Steffe, Eric K.

REFERENCE/DOCKET NUMBER: 1368

REFERENCE/DCCKET NUMBER: 1368

REFERENCE/DCCKET NUMBER: 1360

TELECOMMUNICATION NUMBER: 1360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3 CTACACGCGGGATGCCTGGCTCTTTGGGGCCCTCGTC 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             %; Score 69; DB 3;
; Pred. No. 1.4e+02;
33; Mismatches 0
                                                                                          52241-C/JPW/ADM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 38, Application US/08691814B Patent No. 5981218 GENERAL INFORMATION:
ATTORNEY/AGENT INPORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFRENCE/DOCKET NUMBER: 52241
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-278-0400
TELEPRAK: 212-391-0525
INPORMATION FOR SEQ ID NO: 46:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: not relevant
TOPOLOGY: not relevant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4; Conservative
                                                                                                                                                                                                                                                        LENGTH: 45 amino acids
                                                                                                                                                                                                                                                                                                                   single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; MOLECULE TYPE: peptide US-08-691-814B-38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE:
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: si
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: no
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 4; Conserv
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APPLICANT: Tomasetto, Catherine
APPLICANT: Tomasetto, Catherine
APPLICANT: Basset, Paul
APPLICANT: Byrne, Jennifer
TITLE OF INVENTION: Isolated Nucleic Acid Molecules Useful
TITLE OF INVENTION: as Leukemia Markers and in Breast Cancer Prognosis
NUMBER OF SEQUENCES: 124
CORRESPONDENCE ADDRESS:
STDDEM
STDDEM
                                                Gaps
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Query Match 100.0%; Score 69; DB 2; Length 46; Best Local Similarity 10.8%; Pred. No. 1.4e+02; Matches 4; Conservative 33; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Sterne, Kessler, Goldstein & Fox P.L.L.C.
STREET: 1100 New York Ave, NW, Suite 600
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/691,814B
FILING DATE: 31-JUL-1996
CLASSIFICATION: 435
                                                                                                                       3 CPKCNEFARTSGDHRPCLKCKCGKTSGGHAEHGCHPC 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4 CPKCDTFASSGDHFCLKCRCNKTPGGHAEHDGFCKPC 40
                                                                                          1383.0090001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/002,183
FILING DATE: 09-AUG-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                      Sequence 39, Application US/08691814B Patent No. 5981218 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Steffe, Eric K.
REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 134
TELECOMMUNICATION: TELEPHONE: 202-371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS: not relevant
TOPOLOGY: not relevant
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 47 amino acids
TYPE: amino acid
STRANDEDNESS: not relev
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 202-371-2543
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 10.8
Matches 4; Conservative
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Sequence 183,

US-10-054-611-183

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                                                                                                                                                                                                                          May 27, 2005, 00:35:31 ; Search time 138 Seconds (without alignments) 92.493 Million cell updates/sec
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1: /cgn2_6/ptodata/1/pubpaa/VSO7_PUBCOMB.pep:*
2: /cgn2_6/ptodata/1/pubpaa/PCT7_MEW_PUB.pep:*
3: /cgn2_6/ptodata/1/pubpaa/NSO6_NEW_PUB.pep:*
4: /cgn2_6/ptodata/1/pubpaa/USO6_NEW_PUB.pep:*
5: /cgn2_6/ptodata/1/pubpaa/USO6_NEW_PUB.pep:*
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7: /cgn2_6/ptodata/1/pubpaa/USO8_NEW_PUB.pep:*
8: /cgn2_6/ptodata/1/pubpaa/USO8_NEW_PUB.pep:*
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19: /cgn2_6/ptodata/1/pubpaa/USO0P_PUBCOMB.pep:*
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Copyright (c) 1993 - 2005 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1462099 segs, 344972447 residues
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                             OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 200000000
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69
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Database
                                                                                                                                                                                                                                 Run on:
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Description	Sequence 294, App	Sequence 40, Appl	Sequence 40, Appl	Seguence 46, Appl	Sequence 33022, A	Sequence 8, Appli	Sequence 33565, A	Sequence 8, Appli	Sequence 31, Appl	Sequence 183, App	Sequence 183, App	Sequence 183, App	Sequence 183, App
QI.	US-10-351-334-294	US-08-736-019-40	US-10-844-218-40	US-08-900-230-46	US-10-029-386-33022	US-08-900-230-8	US-09-864-761-33565	US-10-864-774-8	US-10-347-542-31	US-09-843-676-183	US-09-438-486-183	US-10-053-758-183	US-10-054-295-183
DB	15	۵	17	œ	14	ω	σ	17	15	σ	10	14	14
% Query Match Length DB ID	37	39	39	45	47	20	20	20	53	54	54	54	24
% Query Match	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0
Score	69	69	69	69	69	69	69	69	69	69	69	69	69
Result No.	7	8	٣	4	2	9	7	80	σı	10	11	12	13

SUMMARIES

14 69 1000 54 14 US-10-029-186-31820 15 69 1000 0 54 15 US-10-029-186-31820 17 69 1000 0 54 15 US-10-029-186-31820 18 69 1000 0 55 11 US-10-029-186-31820 20 69 1000 0 57 15 US-10-026-1873-1819 21 69 1000 0 57 15 US-10-028-1819992 22 69 1000 0 66 15 US-10-44-599-186942 23 69 1000 0 66 15 US-10-44-599-186942 24 69 1000 0 66 15 US-10-44-599-186942 25 69 1000 0 66 15 US-10-44-599-186942 26 69 1000 0 66 15 US-10-44-599-186942 27 69 1000 0 66 15 US-10-44-599-186942 28 69 1000 0 66 15 US-10-44-599-186942 29 69 1000 0 70 15 US-10-44-599-18642 29 69 1000 0 70 11 US-10-191-391-30 29 69 1000 0 70 11 US-10-101-391-30 20 1000 0 70 11 US-10-101-391-30 20 1000 0 70 11 US-10-101-391-30 20 1000 0 70 11 US-10-269-806-114 20 69 1000 0 70 11 US-10-269-806-114 20 69 1000 0 70 11 US-10-101-391-30 20 1000 0 70 11 US-10-269-806-114 20 1000 0 70 11 US-10-101-391-30 20 1000 0 70 11 US-10-269-806-114 20 1000 0 70 11 US-10-101-391-30 20 1000 0 70 11 US-10-101-391-30 20 1000 0 70 11 US-10-269-806-114 20 1000 0 70 11 US-10-101-391-30 20 1000 0 70 11 US-10-269-806-114 20 1000 0 70 11 US-10-101-391-30 20 1000 0 70 11 US-10-269-806-114 20 1000 0 70 11 US-10-101-391-30 20 1000 0 70 11 US-10-101-391-30 20 1000 0 70 11 US-10-269-806-114 20 1000 0 70 11 US-10-101-391-30 20 1000 0 70 11 US-10-269-806-114 20 1000 0 70 11 US-10-101-391-30 20 1000 0 70 11 US-10-269-806-114 20 1000 0 70 11 US-10-269-806-114 20 1000 0 70 11 US-10-101-391-30 20 1000 0 70 11 US-10-269-806-114 20 1000 0	Sequence 183, App Sequence 183, App Sequence 302, App Sequence 302, App Sequence 4, Appli Sequence 4, Appli Sequence 191604, Sequence 159902, Sequence 159902, Sequence 159902, Sequence 159900, Sequence 159801, Sequence 159801, Sequence 251423, Sequence 251423, Sequence 251423, Sequence 251423, Sequence 15, Appl Sequence 31, Appl Sequence 114, Appl Sequence 114, Appl Sequence 114, Appl Sequence 114, Appl Sequence 115, Appl Sequence 116, Appl Sequence 116, Appl Sequence 117, Appl Sequence 117, Appl Sequence 118, Appl Sequence 15, Appl Sequence 15, Appl		Length 37;
	6 6 9 100.0	RESULT 1 US-10-351-334-294 ; Sequence 294, Application US/10351334 ; Sequence 294, Application US/10351334 ; Publication No. US20040034196A1 ; GENERAL INFORMATION: ; APPLICANT: Komatsoulis et al ; TITLE OF INVENTION: 98 Human Secreted Proteins ; FILE REFERENCE: PZ031P2 ; CURRENT PELING DATE: 2003-01-27 ; PRIOR APPLICATION NUMBER: 06/350, 898 ; PRIOR PELING DATE: 2000-01-25 ; PRIOR PELING DATE: 2000-01-24 ; PRIOR PELING DATE: 1999-07-29 ; PRIOR PILING DATE: 1999-07-29 ; PRIOR PILING DATE: 1998-05 ; PRIOR PELICATION NUMBER: 60/095, 486 ; PRIOR APPLICATION NUMBER: 60/095, 486 ; PRIOR APPLICATION NUMBER: 60/095, 454 ; PRIOR APPLICATION NUMBER: 60/095, 454 ; PRIOR APPLICATION NUMBER: 60/095, 455 ; PRIOR PILING DATE: 1998-08-06 ; PRIOR PILING DATE: 1998-08-06 ; PRIOR PILING DATE: 1998-08-06 ; ROFWARE: PATE PATENTION NUMBER: 60/095, 455 ; PRIOR PILING DATE: 1998-08-06 ; NUMBER OF SEQ ID NOS: 376 ; SEQ ID NO 294 ; LENGTH: 37 ; TYPE: PATE ; TYPE: PATE ; TYPE: PATE ; TYPE: PATE	Match 100.0%; Score 69; DB 15; Scoral Similarity 10.8%; Pred. No. 5.8e+02;

g 2

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Minghetti, Luisa; Waterfield, Michael; Marchioni, Mark; Chen, Maio Su; Hiles, Ian
TITLE OF INVENTION: Glial Mitogenic Factors, Their
Preparation and Use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 10022
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 360 kb storage
                                             Length 39;
                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      | ...|......| ccckccrtcacaaaggtcttctccttcagc 39
                                             100.0%; Score 69; DB 8;
10.8%; Pred. No. 6.2e+02;
ive 33; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 30-JUN-1992
APPLICATION NUMBER: 07/663,703
FILING DATE: 03-APRIL-1992
APPLICATION NUMBER: U.K. 91 07566.3
FILING DATE: 10-APRIL-1991
ATTORNEY/AGENT INFORMATION:
NAME: Teal, Christine H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Tsai, Christine H.
REGISTRATION NUMBER: 34,266
REFERENCE/DOCKET NUMBER: LUD 5250.4
                                                                                                                                                                                                                                                                                                                                                                     Sequence 40, Application US/10844218
Publication No. US20050106666A1
GENERAL INFORMATION:
APPLICANT: Goodearl, Andrew; Stroobant,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: PC-DOS
SOFTWARE: Wordperfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/844,218
FILING DATE: 12-ANG-2004
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/036,555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE DESCRIPTION: SEQ ID NO: 40:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 24-MAR-1993
APPLICATION UNDBER: 07/965,173
FILING DATE: 23-0CT-1992
APPLICATION NUMBER: 07/940,389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 03-SEP-1992
APPLICATION NUMBER: 07/907,138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Felfe & Lynch
STREET: 805 Third Avenue
CITY: New York City
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (212) 838-3884
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (212) 838-:
INFORMATION FOR SEQ ID NO: 40
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 10.8%
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FELEPHONE: (212)
                                                                                                               4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
                                                      Query Match
Best Local Similarity
Matches 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-844-218-40
                                                                                                                                                                                                                                                                                                                     RESULT 3
US-10-844-218-40
US-08-736-019-40
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   Gaps
      0
      Indels
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CORRESPONDENCE ADDRESS:
                                                            1 CDDCQHNTEGQYCQRCKPGFYRDLRRPFSAPDACKPC 37
         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Bieker-Brady, Kristina
REGISTRATION NUMBER: 39,109
REFERENCE/DOCKET NUMBER: 04585/002000
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: 18M Compatible Pentium COMPUTER: 18M Compatible Pentium OPERATING SYSTEM: Windows95 SOFTWARE: FastSeq Version 2.0 CURRENT PEPLICATION DATA: APPLICATION NUMBER: US/08/736,019 FILING DATE: 22-OCT-1996 CLASSIFICATION NUMBER: US/08/736,019 FILING DATE: 06-UN-1995 PRIOR APPLICATION NUMBER: 08/471,833 FILING DATE: 24-DAR-1995 PRIOR APPLICATION NUMBER: 08/936,555 FILING DATE: 23-OCT-1992 PRIOR APPLICATION NUMBER: 07/965,173 FILING DATE: 33-OCT-1992 PRIOR APPLICATION NUMBER: 07/965,173 FILING DATE: 30-UN-1992 PRIOR APPLICATION NUMBER: 07/967,138 FILING DATE: 03-SEP-1992 PRIOR APPLICATION NUMBER: 07/963,703 FILING DATE: 03-APR-1992 PRIOR APPLICATION NUMBER: 07/963,703 FILING DATE: 03-APR-1992 PRIOR APPLICATION NUMBER: US/963,703 FILING DATE: 03-APR-1992 PRIOR APPLICATION NUMBER: US/963,703 FILING DATE: 10-APR-1991 ATTORNEY AGENT INFORMATION DATE: NAME: 07/962,37 PATTORNEY AGENT INFORMATION NAME: 07/962,37 PATTORNEY AGENT AFFACTORY NAME: 07/962,37 PATTORNEY AGENT AFFACTORY NUMBER: US/91 ATTORNEY A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                  Sequence 40, Application US/08736019
Publication No. US20030207799A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     E: Clark & Elbing LLP
176 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                    Stroobant, Paul
Minghetti, Luisa
Waterfield, Michael
Marchionni, Mark
         33;
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(617) 428-7045
                                                                                                                                                                                                                                                                                                                                                            Goodearl, Andrew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chen, Mario
Hiles, Ian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: Massachusetts
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: 176 F
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02110
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 8, Application US/08900230
Publication No. US20010009766A1
GENERAL INFORMATION:
APPLICANT: Bard, Jonathan A.
TITLE OF INVENTION: DNA ENCODING GALANN GALR3 RECEPTORS AND TITLE OF INVENTION: USES THEREOF NUMBER OF SEQUENCES: 59
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                           Length 47;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 50;
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                                          OTHER INFORMATION: MAP TO ACO02053.2

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 6

OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 7.4

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 1.5

OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 5.7

OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 5.7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER FRADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: EN PC Compatible
COMPUTER: EN PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/900,230
FILING DATE: 23-JUL-1997
CLASSIFICATION: 435
ATTONENY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFRENCE/DOCKET NUMBER: 52241-C/JPW/ADM
TELECOMMUNICATION INFORMATION:
TELEPRAS: 212-278-0400
TELEPRAS: 212-278-0400
                                                                                                                                                                                                                                                                                                                                             %; Score 69; DB 14;
; Pred. No. 7.9e+02;
33; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 100.0%; Score 69; DB 8; Best Local Similarity 10.8%; Pred. No. 8.6e+02; Matches 4; Conservative 33; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          E: Cooper & Dunham LLp
1185 Avenue of The Americas
                                                                                                                                                                                                                           100.0%;
10.8%; P
                                                                                                                                                                                                                                                                   4; Conservative
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 50 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS: single
  ORGANISM: Homo sapiens
                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amino acid
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: U.S.A.
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STATE: New York
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                                                                                                                                                                               US-10-029-386-33022
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US-09-864-761-33565
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STREET: 116
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APPLICANT: Rank, David R.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David R.
APPLICANT: HISTOR.
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TITLE OF INVENTION: HUMAN GENOME-DERIVES TWO
FILE REPERENCE: AECMICA-X-2
CURRENT APPLICATION NUMBER: US/10/029,386
CURRENT FILING DATE: 2001-12-20
NUMBER OF SEQ ID NOS: 34288
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                                                                               US-08-900-230-46

Sequence 46, Application US/08900230

Publication No. US2010009766A1

GENERAL INFORMATION:
APPLICANT: Bard, Jonathan A.

TITLE OF INVENTION: DNA ENCODING GALANN GALR3 RECEPTORS AND TITLE OF INVENTION: USES THEREOF
NUMBER OF SEQUENCES: 59

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           %; Score 69; DB 8; Length 45;
; Pred. No. 7.5e+02;
33; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                STALE.

COUNTRY: U.C.
ZIP: 11036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUBER: US/08/900,230
FILING DATE: 23-UU-1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
MANNE: White, John P
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 28,678
FREFERENCE/DOCKET NUMBER: 28,678
TELEPHONE: 212-278-0400
TELEPHONE: 212-39-0400
TELEPHONE: 212-39-0400
TOTALED TOTALES:
TELEPHONE: 212-39-0400
TOTALES:
TELEPHONE: 212-39-0400
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SEQ ID NO 33022
LENGTH: 47
3 CGCCKCCRTTCACRCAGAAGGTCTTCTCCTTCTCAGC 39
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STREET: 1185 Avenue of The Americas
CITY: New York
STATE: New York
COUNTRY: U.S.A.
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Publication No. US20030194704A1 '
GENERAL INFORMATION:
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Best Local Similarity 10.8%;
Matches 4; Conservative 3
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STRANDEDNESS: si
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
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US-10-029-386-33022
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Length 53;
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0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Wilson, David
APPLICANT: Wilson, David
APPLICANT: Wo. US20040009530Alk, Steffen
APPLICANT: Wo. US20040009530Alk, Steffen
APPLICANT: Wo. Jiangchun
APPLICANT: Uemura, Robyn
TITLE OF INVENTION: ENGINERED BINDING PROTEINS
FILE REPERENCE: 11001-006-999
CURRENT APPLICATION NUMBER: US/10/347,542
CURRENT PILING DATE: 2003-01-16
NUMBER OF SEQ ID NOS: 46
SOFTWARE: Patentin version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 69; DB 15;
10.8%; Pred. No. 9.3e+02;
tive 33; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             h 100.0%; Score 69; DB 17; Similarity 10.8%; Pred. No. 8.6e+02; 4; Conservative 33; Mismatches 0;
                                                                                                                                                                                                                          Sequence 8, Application US/10864774

Publication No. US2005009646A1

GENERAL INFORMATION:
APPLICANT: SULLIVAN, SEAN M.
TITLE OF INVENTION: GENE DELIVERY TO TUMORS
FILE REFERENCE: 5853-337-1WO
CURRENT APPLICATION NUMBER: US/10/864,774

CURRENT FILING DATE: 2004-06-09

NUMBER OF SEQ ID NOS: 18

SOFTWARE: Patentin version 3.2
  Mismatches
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Sequence 183, Application US/09843676
Patent No. US20020164786A1
GENERAL INFORMATION:
APPLICANT: Cech, Thomas R.
Lingner, Joachim
Nakamura, Toru
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Sequence 31, Application US/10347542; Publication No. US20040009530A1; GENERAL INFORMATION:
  33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Pyrococcus furiosus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         , ORGANISM: Rattus norvegicus
US-10-864-774-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
  Conservative
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
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US-09-843-676-183
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        4,
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US-10-864-774-8
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LENGTH: 50
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LENGTH: 53
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                                                                                 APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David K.
APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
FILE REFERENCE: Aeomica-X-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NN: MAP TO AP000091.1

NN: EXPRESSED IN BONE MARROW, SIGNAL = 9

NN: EXPRESSED IN LUNG, SIGNAL = 12

NN: EXPRESSED IN PLACENTA, SIGNAL = 12

NN: EXPRESSED IN PLACENTA, SIGNAL = 7.5

NN: EXPRESSED IN HEL10, SIGNAL = 9

NN: EXPRESSED IN HELA, SIGNAL = 9

NN: EXPRESSED IN HELA, SIGNAL = 12

NN: EXPRESSED IN HELA, SIGNAL = 8.7

NN: EXPRESSED IN HELA, SIGNAL = 15

NN: EXPRESSED IN HEART, SIGNAL = 15

NN: EXPRESSED IN HEART, SIGNAL = 15

NN: EXPRESSED IN HEART, SIGNAL = 11

NN: EXPRESSED IN HEART SIGNAL = 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Annomax Sequence Listing Engine vers. 1.1 SEQ ID NO 33565 LENGTH: 50
                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
PRIOR PILLING DATE: 2000-02-04
PRIOR FILING DATE: 2000-02-04
PRIOR FILING DATE: 2000-05-26
PRIOR PILLING DATE: 2000-05-26
PRIOR PILLING DATE: 2000-06-03
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR PILLING DATE: 2000-00-03
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR PILLING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 09/174,203 PRIOR FILING DATE: 2001-01-29
        Sequence 33565, Application US/09864761 Patent No. US20020048763A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 49117
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: E OTHER INFORMATION: E
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Best Local Similarity
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OTHER INFORMATION:
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US-09-864-761-33565
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Best Local Similarity 10.8%; Pred. No. 9.5e+02;
Matches 4; Conservative 33; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Harley, Caĭvin
Andrews, William H.
TITLE OF INVENTION: No. US20030032075Alel Telomerase
NUMBER OF SEQUENCES: 225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, 8th Floor CITY: San Francisco
                                                                              ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/438,486
FILING DATE: 12-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3 CSACSTISGRGAPASWAPLCWAWTISIGPGAPSCCVC 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION UNDRER:
APPLICATION UNDRER:
FILING DATE: 01-0CT-1996
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: APPLE REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002931US
Two Embarcadero Center, 8th Floor
                                                                                                                                                                                                                                                                                      CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
CLASSIFICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
CLASSIFICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
PRIOR PAPLICATION DATA:
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
FILING DATE: 18-APR-1997
                                                                  United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 183, Application US/10053758; Publication No. US20030032075A1 GENERAL INFORMATION: APPLICANT: Cech, Thomas R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nakamura, Toru
Chapman, Karen B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lingner, Joachim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Morin, Gregg B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FELECOMMUNICATION INFORMATION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        54 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide
                     San Francisco
California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE:
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                                          STATE: Ca
COUNTRY:
ZIP: 9411
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                                                                Andrews, William H.
TITLE OF INVENTION: No. US20020164786Alel Telomerase
NUMBER OF SEQUENCES: 225
                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/843,676
FILING DATE: 26-Apr-2001
CLASSIFICATION: 536
                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSER: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, 8th Floor CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Andrews, William H.
TITLE OF INVENTION: No. US20030009019A1el Telomerase
NUMBER OF SQUENCES: 223
CORRESPONDENCE ADDRESS:
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REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002930US
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APPLICATION NUMBER: US/08/854,050
FILING DATE: 09-MAY-1997
PRILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-OCT-1996
                                                                                                                                                                                                                                              COUNTRY: United States of America ZIP: 94111
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 183:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 183, Application US/09438486
Publication No. US2003000901941
GENERAL INFORMATION
APPLICANT: Cech, Thomas R.
APPLICANT: Lingner, Joachim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (415) 576-0200
(415) 576-0300
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  Chapman, Karen B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 54 amino acids
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Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Hailey, Calvin
                          Morin, Gregg B.
Harley, Calvin
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Best Local Similarity 10.8*
Matches 4; Conservative
                                                                                                                                                                                                                             STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-438-486-183
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Gaps
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TITLE OF INVENTION: No. US20030059787A1el Telomerase
NUMBER OF SEQUENCES: 225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew STREET: Two Embarcadero Center, 8th Floor CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                    NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REPREBNCE/DOCKET NUMBER: 015389-002930US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3 CSACSTTSGRGAPASWAPLCWAWTISTGPGAPSCCVC 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
100.0%; Score 69; DB 14;
Best Local Similarity 10.8%; Pred, No. 9.5e+02;
Matches 4; Conservative 33; Mismatches 0:
                                                                                                                          APPLICATION DATE: 08/854,050
FILING DATE: «Unknown»
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-OCT-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: California
COUNTRY: United States of America
ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/054,295
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/054,611
FILING DATE: 18-Jan-2002
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 183:
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Publication No. US20030059787A1
GENERAL INFORMATION:
APPLICANT: Cech, Thomas R.
Lingner, Joachim
                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (415) 576-0200
                                                               FILING DATE: 18-Jan-2002
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                   (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDEDNESS: <Unknown>
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Chapman, Karen B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 54 amino acids
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                                                                                                              PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ropology: linear
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; Pred. No. 9.5e+02;
33; Mismatches 0; Indels
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Harley, Calvin
Andrews, William H.
TITLE OF INVENTION: NO. US20030044953Alel Telomerase
NUMBER OF SEQUENCES: 225
CORRESPONDENCE ADDRESS:
                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, 8th Floor TTTY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Apple, Randolph T.
REGISTRADIN NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002930US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/854,050 FILING DATE: 09-M2x-1997 APPLICATION NUMBER: US 08/851,843 APPLICATION NUMBER: US 08/846,017 APPLICATION NUMBER: US 08/846,017 APPLICATION NUMBER: US 08/844,419 FILING DATE: 18-APR-1997 FILING DATE: 18-APR-1997 FILING DATE: 01-007-1996
                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/053,758
FILING DATE: 18-Jan-2002
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
  United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 183:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 183, Application US/10054295
Publication No. US20030044953A1
GENERAL INFORMATION:
APPLICANT: Cech, Thomas R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: <Unknown>
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Nakamura, Toru
Chapman, Karen B.
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10.8%; E
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                         ZIP: 94111
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4; Conservative
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Best Local Similarity
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US-10-054-295-183
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Sequence 33820 Application US/10029386
Sequence 33820 Application US/10029386
Publication No. US20030194704A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Penn, Sharron G.
APPLICANT: Hanzel, David R.
APPLICANT: Hanzel, David R.
APPLICANT: HANSHION: EXPRESSION ANALYSIS TWO
TITLE OF INVENTION: EXPRESSION ANALYSIS TWO
FILE REFERENCE: AEOMICA-X-2
CURRENT APPLICATION NUMBER: US/10/029,386
CURRENT FILING DATE: 2001-12-20
NUMBER OF SEQ ID NOS: 34288
SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
SEQ ID NO 33820
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 69; DB 14; Length 54;
Pred. No. 9.5e+02;
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CTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 1.8
CTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 1.9
CTHER INFORMATION: SWISSPROT HIT: P34088, EVALUE 1.60e+00
US-10-029-386-33820
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: ADD1e, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002930US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3 CSACSTTSGRGAPASWAPLCWAWTISTGPGAPSCCVC 39
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FILING DATE: «Unknown»
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
FILING DATE: 01-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              33; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear

MOLECULE TYPE: peptide

SEQUENCE DESCRIPTION: SEQ ID NO: 183:
US-10-054-611-183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Search completed: May 27, 2005, 00:48:47 Job time : 149 secs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 183:
SEQUENCE CHARACTERISTICS:
LENGTH: 54 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
10.8%; P
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Best Local Similarity 10.8%
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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US-10-029-386-33820
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00:26:25; Search time 39 Seconds (without alignments) 91.283 Million cell updates/sec
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Copyright (c) 1993 - 2005 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                            283416 segs, 96216763 residues
                                                                        using sw model
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Maximum DB seq length: 200000000
                                                                                                                                                                              US-09-763-616-2
69
                                                                                                        May 27, 2005,
                                                                    OM protein .- protein search,
                                                                                                                                                                                                                                                     BLOSUM62DX
                                                                                                                                                                                                   Perfect score:
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                                                                                                                                                                                                                                                                                                            Searched:
                                                                                                        Run on:
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Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries pir2:* pir3:* pir4:* PIR 79:* Database

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	rubredoxin [valida	rubredoxin - Desul	rubredoxin (valida	rubredoxin [valida	rubredoxin - Helio	probable rubredoxi	rubredoxin - Chlor	rubredoxin - Clost	rubredoxin - "Buty	ä	rubredoxin - Therm	rubredoxin - Metha	rubredoxin PAB7224	rubredoxin - Clost	rubredoxin [valida	rubredoxin [valida	rubredoxin [import	rubredoxin - Acine	rubredoxin [import	probable rubA prot	rubredoxin PA5350	. rubredoxin PA5351	testis-specific pr		'n	ferredoxin (fdx-8)	rubredoxin XF0379	ferredoxin [import	ferredoxin 2[4Fe-4
SUMMARIES	ID	A33173	JX0241	RUDV	RUDVEG	S65620	G71256	A27537	A33182	JU0127	H69359	H72348	B69075	E75136	JU0074	RUCLEP	RUPF	G97241	139520	AE2375	E70593	H82976	A82977	S25774	WTFF	H81133	F69517	B82814	B97333	B42960
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de	Query Match	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0
	Score	69	69	69						69				69				69		69						69	69		69	69
	Result No.	-	7	m	4	S	9	7	œ	6	10	11	12	13	14		16	17	18	19	20	21		23	24	25	56	27	28	29

ferredoxin 2[4Fe-4	rubredoxin [import	probable rubB prot	ferredoxin (fdx-6)	rubredoxin - Metha	testis-specific pr	hemagglutinin VCAO	ferredoxin - Therm	testis-specific pr	hupl protein - Rhi	hydrogenase-relate	rubredoxin homolog	rubredoxin (rd-2)	testis-specific pr	Mid-ORF2 protein -	anaredoxin [import
FEDV2N	F87558	D70593	B69376	G69073	S25772	S18996	A72324	S25775	S27340	853661	E44915	D69418	\$25773	D36760	AB1988
7	7	~	,7	7	7	7	7	~	N	0	7	7	~	~	0
29	59	9	9	63	63	63	65	68	70	72	72	73	74	74	16
100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0
69	69	69	69	69	69	69	69	69	69	69	69	69	69	69	69
30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45

ALIGNMENTS

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Tubredoxin [validated] - Clostridium thermosaccharolyticum (Species: Clostridium thermosaccharolyticum, Clostridium tartarivorum (Species: Clostridium thermosaccharolyticum, Clostridium tartarivorum (Spacies: 30-Apr-1991 #sequence_revision 24-Oct-1997 #text_change 09-Jul-2004 (Spacession: A33173 R.Meyer, J.; Gagnon, J.; Sieker, L.C.; Van Dorsselaer, A.; Moulis, J.M. Babedoxin, 1.371, 1990 A;Title: Rubredoxin from Clostridium thermosaccharolyticum. Amino acid sequence, mass-s A;Reference number: A33173; MUID:91058526; PMID:2244884
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tubredoxin - Desulfovibrio vulgaris (strain Miyazaki)
C;Species: Desulfovibrio vulgaris
C;Species: Desulfovibrio vulgaris
C;Species: 10-Un-1993 #sequence_revision 22-Unl-1994 #text_change 09-Jul-2004
C;Accession: JX0241
R;Shimizu, F.; Ogata, M.; Yaqi, T.; Wakabayashi, S.; Matsubara, H.
B;Chimizu, F.; Ogata, M.; Yaqi, T.; Wakabayashi, S.; Matsubara, H.
A;Chimizu, F.; Ogata, M.; Yaqi, T.; Wakabayashi, S.; Matsubara, H.
A;Atitle: Amino acid sequence and function of rubredoxin from Desulfovibrio vulgaris Miy
A;Reference number: JX0241; WUID:90234754; PMID:2561345
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C,Superfamily: rubredoxin; rubredoxin homology
C,Keywords: electron transfer; iron; metalloprotein
F3.349/Domain: rubredoxin homology (Wpartial) #status experimental
F;1/Modified site: N-formylmethionine (partial) #status experimental
F;6,9,39,42/Binding site: iron (Cys) #status predicted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                A.Molecule type: protein
A.Residues: 1-52 -MEY>
A.Cross-references: UNIPROT:P19500
C.Superfamily: rubredoxin homology
C.Keywords: electron transfer; iron; metalloprotein
F.3-49/Domain: rubredoxin homology RUB>
F.3-49/Domain: rubredoxin homology RUB>
F.3-49/Damain: rubredoxin homology RUB>
F.3-49/Damain: rubredoxin homology RUB>
F.3-49/Damain: rubredoxin homology RUB>
F.3-49/Binding site: iron (Cys) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |::|::::::::|:::|CTVCGYIYDPEVGDPIQNIPPGTKFEDLFDDWVCPDC 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 69; DB 1;
10.8%; Pred. No. 3.1e+02;
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33; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity 10.8%; 4; Conservative 3
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A;Residues: 1-52 <SHI>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 4; Conserv
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Best Local Similarity
Matches 4; Conserv
                                                                                                                                                                                                                                                                                                         A; Accession: A33173
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Gaps

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C;Accession: S65620
R;Lee, W.Y.; Brune, D.C.; LoBrutto, R.; Blankenship, R.E.
Arch. Blochem. Biophys. 318, 80-88, 1995
A;Title: Isolation, characterization, and primary structure of rubredoxin from the phot A;Reference number: 865620; MUID:95243660; PMID:7726577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: UNIPROT:O83956; GB:AE001266; GB:AE000520; NID:g3323309; PIDN:AAC65:
A;Experimental source: strain Nichols
C;Genetics:
                                                                                                                                                                                                           of the oxidized form at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A, Residues: 1-52 < LEB.
A, Residues: 1-52 < LEB.
A, Residues: 1-52 < LEB.
A, Crose-references: WIPPOT: P56263
C, Crose-references: WIPPOT: p56263
C, Comment: Rubredoxin is a nonheme iron protein and substitutes for ferredoxin in some C, Superfamily: rubredoxin; rubredoxin homology
C, Reywords: electron transfer; hydrocarbon hydroxylation; iron; metalloprotein P; 3-49/Domain: rubredoxin homology < RUB.
F; 1/Modified site: N-formylmethionine #status experimental
F; 6, 9, 39, 42/Binding site: iron (Cys) #status predicted
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Clacession: G71256

Clacession: G71256

Clacession: G.M.; Norris, S.J.; Weinstock, G.M.; White, O.; Sutton, G.G.; Dodson, rson, J.; Khalak, H.; Richardson, D.; Howell, J.K.; Chidambaram, M.; Utterback, they, L.; Weidman, J.; Smith, H.O.; Venter, J.C.

Science 281, 375-388, 1998

A;Title: Complete genome sequence of Treponema pallidum, the syphilis spirochete A;Reference number: A71250; MUID:98332770; PMID:9665876
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Status: preliminary; nucleic acid sequence not shown; translation not shown
submitted to the Brookhaven Protein Data Bank, March 1988
A;Reference number: A50332; PDB:1RDG
A;Contents: annotation; X-ray crystallography, 1.4 angstroms, residues 1-52
R;Frey, M.; Sieker, L.; Payan, F.; Haser, R.; Bruschi, M.; Pepe, G.; LeGall,
J. Mol. Biol. 197, 525-54, 1987
A;Title: Rubredoxin from Desulfovibrio gigas. A molecular model of the oxidiz
A;Reference number: A44624; WUID:88155649; PMID:3441010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   probable rubredoxin - syphilis spirochete
C;Species: Treponema pallidum subsp. pallidum (syphilis spirochete)
C;Date: 24-Jul-1998 #sequence_revision 24-Jul-1998 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  rubredoxin - Heliobacillus mobilis
C;Species: Heliobacillus mobilis
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 09-Jul-2004
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                                                                                                                                                                                                                                                                                               A Contents: annotation; X-ray crystallography, 1.4 angstroms C; Superfamily: rubredoxin, rubredoxin homology C;Keywords: blocked amino end; electron transfer; iron; metalloprotein F;3-49/Domain: rubredoxin homology <RUB>
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10.8%; Pred. No. 3.1e+02;
iive 33; Mismatches 0;
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Pred. No. 3.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  h 100.0%; Score 69; DB Similarity 10.8%; Pred. No. 3.1e-4; Conservative 33; Mismatches
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Best Local Similarity
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A; Residues: 1-52 <COL>
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C;Species: Desulfovibrio gigas
C;Species: Desulfovibrio gigas
C;Date: 07-May-1981 #sequence_revision 07-May-1981 #text_change 09-Jul-2004
C;Accession: A00275
R;Bruschi, M.
Biochem. Biophys. Res. Commun. 70, 615-621; 1976
A;Title: The amino acid sequence of rubredaxin from the sulfate reducing bacterium, Desu
A;Reference number: A00275; MUID:76231572; PMID:938515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bruschi, M.
Jochim. Blophys. Acta 434, 4-17, 1976
Jiochim. Blophys. Acta 434, 4-17, 1976
Jittle: Non-heme iron proteins. The amino acid sequence of rubredoxin from Desulfovibri
Jittle: Non-heme iron proteins. The amino acid sequence of rubredoxin from Desulfovibri
JReference number: A00274; MUID:76232334; PMID:7308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C;Accession: B33962; A00274 —
R;Brumlik, M.J.; Voordouw, G.
B. Bacteriol. 171, 4996-5004, 1989
A;Title: C, the transcriptional unit encoding the genes for rubredoxin (rub) an A;Reference number: A33962; MUID:89359139; PMID:2549009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cross-references: UNIPROT: P00269; GB:M28848; NID:g342028; PIDN:AAA64798.1; PID:g758677
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: protein
A; Residues: 1-20, T', 22-52 <a href="Molecule">Molecule type: protein</a>
A; Residues: 1-20, T', 22-52 <a href="Molecule">Molecule type: protein</a>
A; Residues: 1-20, T', 22-52 <a href="Molecule">Molecule type: protein</a>
B; Experimental source: strain Hildenborough
B; Adman, E.T.; Sieker, L.C.; Jensen, L.H.
A; Reference nundber: A50939; PBS:7RXN
A; Reference nundber: A50939; PBS:7RXN
A; A; Mol. Biol. 217, 337-352, 1991
A; Title: Structure of rubredoxin from Desulfovibrio vulgaris at 1.5 A resolution. A; Reference number: A22814; MUID: 91124457; PMID:1992166
A; Contents: annotation; X ray crystallography, 1.5 angstroms
B; Dauter, Z.; Sieker, L.; Wilson, K.
Bubmitted to the Brookhaven Protein Data Bank, August 1991
A; Reference number: A51562; PBB:RXN
A; Reference number: A51562; PBB:RX
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(Species: Desulfovibrio vulgaris
A;Variety: strain Hildenborough
C;Date: 24-Apr-1984 #sequence_revision 30-Jun-1991 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.0 angstroms, residues 1-52
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                                                                                                                                                                                                                                                                                                                                              rubredoxin [validated] - Desulfovibrio vulgaris (strain Hildenborough)
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Keywords: electron transfer; iron; metalloprotein
F;3-49/Domain: rubredoxin homology «KUB»
F;6,9,39,42/Binding site: iron (Cys) #status experimental
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                                                                   33; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Contents: annotation; X-ray crystallography,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10.8%;
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Best Local Similarity
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C;Species: "Butyribacterium methylotrophicum"
C;Species: "Butyribacterium methylotrophicum"
C;Date: 31-Mar-1990 #sequence_revision 22-Jul-1994 #text_change 09-Jul-2004
C;Date: 31-Mar-1990 #sequence_revision 22-Jul-1994 #text_change 09-Jul-2004
C;Date: 31-Mar-1990 #sequence_revision 22-Jul-1994 #text_change 09-Jul-2004
R;Saeki, K.; Yao, Y.; Wakabayashi, S.; Shen, G.J.; Zeikus, J.G.; Matsubara, H.
J. Biochem. 106, 656-662, 1989
J. Biochem. 106, 656-662, 1989
A;Title: Ferredoxin and rubredoxin from Butyribacterium methylotrophicum: complete prim A;Reference number: A91913; MUID:90110065; PMID:2606914
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tubredoxin (rd-1) homolog - Archaeoglobus fulgidus
C; Species: Archaeoglobus fulgidus
C; Species: Archaeoglobus fulgidus
C; Species: Archaeoglobus fulgidus
C; Species: O5-Dec-1997 #sequence_revision O5-Dec-1997 #text_change O9-Jul-2004
C; Accession: H69359
R; Klenk, H.P.; Clayton, R.A.; Tomb, J.F.; White, O.; Nelson, K.E.; Ketchum, K.A.; Dodso.; Fleischmann, R.D.; Quackenbush, J.; Lee, N.H.; Sutton, G.G.; Gill, S.; Kirkness, B.F. Glodek, A.; Zhou, L.; Overbeek, R.; Gocayne, J.D.; Weidman, J.F.; McDonald, L.
Nature 390, 364-370, 1997
A; Authors: Utterback, T.; Cotton, M.D.; Spriggs, T.; Artiach, P.; Kaine, B.P.; Sykes, S. A; Authors: Utterback, T.; Cotton, M.D.; Spriggs, T.; Artiach, P.; Kaine, B.P.; Sykes, S. A; Title: The complete genome sequence of the hyperthermophilic, sulfate-reducing archae A; Reference number: A69250; MUID:98049343; PMID:9389475
A; Accession: H69359
A; Accession: H69359
A; Accession: A; Mosidues: Drawn, rucleic acid sequence not shown; translation not shown
A; Mosidues: 1-53 «KLE»
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C;Date: 11-Jun-1999 #sequence_revision 11-Jun-1999 #text_change 09-Jul-2004
C;Accession: H72349
R;Nelson, K.E.; Clayton, R.A.; Gill, S.R.; Gwinn, M.L.; Dodson, R.J.; Haft, D.H.; Hicke
Garrett, M.M.; Stewart, A.M.; Cotton, M.D.; Pratt, M.S.; Phillips, C.A.; Richardson, D.
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                                                                                                                                                                                                                                                                                                                                                                     A;Cross-references: UNIPROT:P14071
C;Comment: Rubredoxin is a nonheme iron protein and substitutes for ferredoxin in some
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C;Keywords: electron transfer; iron; metalloprotein
F;F.3-49,0main: rubredoxin homology <RUB>
F;6,9,39,42/Binding site: iron (Cys) #status predicted
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F;6,9,39,42/Binding site: iron (Cys) #status predicted
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Best Local Similarity
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Matches
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Eur. J. Biochem. 163, 161-166, 1987

File: The complete amino acid sequence of rubredoxin from the green phototrophic bact

A; Reference number: A27537; WUID:87133563; PMID:3816795

A; Note: Chlorobium thiosulphatophilum

A; Accession: A27537

A; Addecule type: protein

A; Residues: 1-53 < WOO>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           rubredoxin - Chlorobium limicola f.sp. thiosulfatophilum
C;Species: Chlorobium limicola f.sp. thiosulfatophilum
C;Date: 05-Jun-1988 #sequence_revision 22-Jul-1994 #text_change 09-Jul-2004
C;Accession: A27537
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;1-Modified site: N-formylmethionine #status experimental
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C;Superfamily: rubredoxin; rubredoxin homology
C;Keywords: electron transfer; iron; metalloprotein
F;3-49/Domain: rubredoxin homology <RUB>
F;6,9,39,42/Binding site: iron (Cy8) #status predicted
                      C; Keywords: iron, metalloprotein
P;3-49/Domain: rubredoxin homology <RUB>
F;6,9,39,42/Binding site: iron (Cys) #status predicted
                                                                                                                                                                                                                                                                                                              | ::| ::::::::| | CSVCGYVYDPADGEPDDPIDPGTGFEDLPEDWVCPVC 42
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R;Meyer, J.; Gagnon, J.; Moulis, J.M.
submitted to the Protein Sequence Database, April 1991
A;Reference number: A33182
A;Accession: A33182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CTVCGYVYDPEVGDPDNNINPGTSFQDIPEDWVCPLC 42
                                                                                                                                                                                                                                                                       F;6,9,39,42/Binding site: iron (Cys) #status predicted
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10.8%; Pred. No. 3.1e+02;
ive 33; Mismatches 0
                                                                                                                                               100.0%; Score 69; DB 2;
10.8%; Pred. No. 3.1e+02;
ive 33; Mismatches 0
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10.8%; Pred. No. 3.1e+02;
Superfamily: rubredoxin; rubredoxin homology:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      rubredoxin - Clostridium sticklandii
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A;Residues: 1-53 MBY.
A;Cross-references: UNIPROT:P23474
A;Experimental source: ATCC 12662
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Species: Clostridium sticklandii
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                                                                                                                                                                                                             4; Conservative
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Matches 4; Conserv
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Gaps

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A,Title: Evidence for lateral gene transfer between Archaea and Bacteria from genome se. A,Reference number: A72200; MUID:99287316; PMID:10360571

RESULT

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Gaps

Gaps

0;

Length 53;

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rubredoxin - Clostridium perfringens
C.Species: Clostridium perfringens
C.Species: 28-Feb-1990 #sequence_revision 22-Jul-1994 #text_change 09-Jul-2004
C.Accession: U00074
R.Seki, Y.; Seki, S.; Satoh, M.; Ikeda, A.; Ishimoto, M.
B. Biochem: 106, 336-341, 1989
A.Fitle: Rubredoxin from Clostridium perfringens: complete amino acid sequence and part
A.Reference number: JU0074; MUID:90036784; PMID:2553684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Molecule type: protein
A; Residues: 1-54 <SEK.
A; Cross-references: UNIPROT: P14072
C; Comment: The protein is reduced with NADH in the presence of a specific NAD(P) H oxido
C; Superfamily: rubredoxin; rubredoxin homology
C; Keywords: electron transfer; iron; metalloprotein
F; 3-49, Domain: rubredoxin homology <RUB.
F; 6,9,39,42/Binding site: iron (Cys) #status predicted
                                                                                                                                                                                                               %; Score 69; DB 2; Length 53
; Pred. No. 3.1e+02;
33; Mismatches 0; Indels
A;Gene: rd; PAB7224
C;Superfamily: rubredoxin; rubredoxin homology
C;Reywords: iron; metalloprotein
F;3-49(Domain: rubredoxin homology <RUB>
F;6,9,39,42/Binding site: iron (Cys) #status predicted
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                                                                                                                                                                                                                                                                                                                                                            h Similarity 10.0%; Score 69; DB 1; Similarity 10.8%; Pred. No. 3.1e+02; 4; Conservative 33; Mismatches 0
                                                                                                                                                                                                                   100.0%; Score 69;
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Best Local Similarity 10.8%
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C;Date: 05-Dec-1997 #sequence_revision 05-Dec-1997 #text_change 09-Jul-2004
C;Date: 05-Dec-1997 #sequence R; Vicaire, R; Vicai
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A;Experimental source: strain Orsay
C;Genetics:
                                                                                                                                            :Cross-references: UNIPROT:Q9WZC7; GB:AE001739; GB:AE000512; NID:g4981176; PIDN:AAD3574
:Experimental source: strain MSB8
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R;anonymous, Genoscope
submitted to the EMBL Data Library, July 1999
A;Description: Pyroccoccus abyssi genome sequence: insights into archaeal chromosome A;Reference number: A75001
A;Reference number: E75136
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C;Species: Methanobacterium thermoautotrophicum
C;Date: 05-Dec-1997 #sequence_revision 05-Dec-1997 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              #text_change 09-Jul-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 53;
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                                                                                                                                     A.Cross-references: UNIPROT:09WZC7; GB:AE001739; GB:AE0
A.Experimental source: strain MSB8
G.Genetics:
A.Gene: TM0659
C.Superfamily: rubredoxin; rubredoxin homology
C.Keywords: iron, metalloprotein
F.3-49/Domain: rubredoxin homology «RUB»
F.6-9,39,42/Binding site: iron (Cy8) #status predicted
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C,Superfamily: rubredoxin; rubredoxin homology
C,Keywords: iron; metalloprotein
F;3-49/Domain: rubredoxin homology <RUB>
F;6,9,39,42/Binding site: iron (Cys) #status predicted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 69; DB 2; L
10.8%; Pred. No. 3.1e+02;
ive 33; Mismatches 0;
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10.8%; Pred. No. 3.1e+02;
ive 33; Mismatches 0;
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C;Date: 20-Aug-1999 #sequence_revision 20-Aug-1999
C;Accession: E75136
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Residues: 1-53 <KAW>
                                     A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-53 <ARN>
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   A; Accession: H72348
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0; Indels

Length 54;

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rubredoxin [validated] - Clostridium pasteurianum
C;Species: Clostridium pasteurianum
C;Accession: S29120; A00273
R;Mathieu, I.; Meyer, J.; Moulis, J.M.
B;Jochem. J. 285, 285-286, 1992
B;Jochem. J. 285, 285-286, 1992
B;Jochem. J. 285, 285-286, 1992
B;Jochem. J. 285, 285-281, 1992
B;Jochem. J. 285, 285-281, 1993
B;Jochem. J. 285, 285, 1993
B;Jochem. J. 285, 1993
B;Jochem. J. 285, 285, 1993
B;Joch
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A; Molecule type: protein
A; Residues: 1-13, 'D', 15-21, 'D', 23-47, 'E', 49-54 < MCC>
A; Note: peptides for which chemical overlaps were not determined were positioned on the R; Dauter, Z.; Wilson, K.S.; Sieker, L.C.; Moulis, J.M.; Meyer, J.
Submitted to the Brookhaven Protein Data Bank, December 1995
A; Reference number: A65897; PBI-IRO
A; Reference number: A65897; PBI-IRO
A; Contents: annotation; X-ray crystallography, 1.1 angstroms, residues 1-53
R; Dauter, Z.; Wilson, K.S.; Sieker, L.C.; Moulis, J.M.; Meyer, J.
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submitted to the Brookhaven Protein Data Bank, December 1995
A;Reference number: A65886; PDB:1IRN
A;Contents: annotation; X-ray crystallography, 1.2 angstroms, residues 1-53
R;Watenpaugh, K.D.; Sieker, L.C.; Jensen, L.H.
J. Mol. Biol. 138, 615-633, 1980
A;Title: Crystallographic refinement of rubredoxin at 1.2 Angstroms resolution.
A;Reference number: A56640; MUID:81005589; PMID:7411618
A;Contents: annotation; X-ray crystallography, 1.2 angstroms
R;Watenpaugh, K.D.; Siecker, L.C.; Herriott, J.R.; Jensen, L.H.
A;Reference number: A5009
A;Title: Refinement of the model of a protein: rubredoxin at 1.5 angstrom resolution.
A;Title: Refinement of the model of a protein: rubredoxin at 1.5 angstroms
C;Superfamily: rubredoxin; rubredoxin homology
C;Keywords: annotation; X-ray crystallography, 1.5 angstroms
C;Superfamily: rubredoxin; rubredoxin homology
C;Keywords: electron transfer; fron; metalloprotein
F;1,Modified site: N-formylmethionine #status experimental
F;6,9,39,42/Binding site: iron (Cys) #status experimental
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clostridium clostridium

Q9a194 P00268

acinetobact

acinetobact

anabaena sp chromobacte

bordetella bordetella bordetella

bacteroides

mycobacteri

drosophila drosophila drosophila

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MEDIATES PROM N.A.

MEDIATRES POSTS 57; PubMed=2510401;

MEDIATRES POSTS 57; PubMed=2510401;

MARCHINES POSTS 57; PubMed=2510401;

MARCHINES POSTS 58; Manaye E.M.;

The completed sequence of lymphocytic choriomeningitis virus reveals a unique RNA structure and a gene for a zinc finger protein.";

Virology 173:1-10(1989).

LI VIROLOGY 173:1-10(1989).

MIGHERPO, IPRO07087; Zaf CRH2.

RICHERPO, IPRO07087; Zaf CRH2.

RICHERPO, IPRO07087; Zaf P11.

RR Pfam; PF03654; Zaf P11; 1.

RR PF050m; PD020016; Znf P11; 1.

RR PROSITE; PS00028; ZINC_FINGER_CCH2_1; 1.

MARA-binding; Zinc_finger.

TINON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                             Lymphocytic choriomeningitis virus (strain Pasteur).
Viruses; ssRNA negative-strand viruses; Arenaviridae; Arenavirus.
VCBI_TaxID=11625;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Name=TubA; OrderedLocusNames=XCC3268;
Xanthomonas campestris (pv. campestris).
Bacteria; Proteobacteria; Gammaproteobacteria; Xanthomonadaceae; Xanthomonas.
NCBI_TaxID=340;
                                            09aqjs
08ynk8
07plr1
07w047
07wf74
08a4q8
001644
06qar9
005894
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              h Similarity 10.0%; Score 69; DB 1; Length 51; Similarity 10.8%; Pred. No. 5.16+02; 4; Conservative 33; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5972 MW; 1B2E383153CD8582 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11 CKSCWQKFDSLVRCHDHYLCRHCLNLLLTSSDRCPLC 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-NOV-1990 (Rel. 16, Created)
01-NOV-1990 (Rel. 16, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
                                                                                                                                                                                                                                                     ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                    51
RUBR_ACIAD
BURBR_CLOPA
COAB
RUBR_CLOPA
COAQUINKB
COPYNKB
COPYNKB
COTWAITA
COWNOTA
COWNOTA
COWNOTA
COWNOTA
COMMOTA
COMMOTA
COMMOTA
COMMOTA
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COMMOTA
COMMOTA
COMMOTA
COACA
CO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-OCT-2002 (TrEMBLrel. 22, 01-OCT-2003 (TrEMBLrel. 25, Rubredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-OCT-2002 (TrEMBLrel. 22,
                                                                                                                                                                                                                                                                                                                                    STANDARD;
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 Local Similarity
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 1000.0
1000.0
1000.0
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P19326;
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SEQUENCE
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Matches
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desulfovibr
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clostridium
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bacteroides
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                                                                                                     May 27, 2005, 00:25:40 ; Search time 175 Seconds (without alignments) 108.268 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Description
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Q8ph46 3
P58993 0
P19500 0
P00270 0
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P56263
Q083956
Q08709
Q074709
Q74700
Q74700
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Q74707
P14072
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P24297
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Q46496
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Compugen Ltd.
                                                                                                                                                                                                                                                                                                              Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                 1612378 segs, 512079187 residues
              GenCore version
Copyright (c) 1993 - 2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZNFP LYCVP
Q8P5R8
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RUBR_CLOTS
RUBR_DESGI
RUBR_DESVH
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RUBR_HELMO
RUBR_TREPA
Q6QAŪ0
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RUBR_BUTME
RUBR_CHLLT
RUBR_CLOST
RUBR_PYRAB
RUBR_PYRFU
026259
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Maximum Match 100%
Listing first 45 summaries
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RUB2 CLOPE
                                                                           protein search, using sw model
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Q8R870
Q73NV0
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Q74EW2
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Q9FDN9
Q64N49
Q9WZC7
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1: uniprot_sprot:*
2: uniprot_trembl:*
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length: 2000000000
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69
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seq

Minimum DB Maximum DB

Database

Result No.

Scoring table:

Searched:

Perfect score:

Sequence:

OM protein -

Run on:

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Gaps

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InterPro; IPR004039; Rubredox.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                         CHLTE
                                                                                                                                                                                                                     Complete | SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                          RUB2_CHLTE
ID RUB2_CHL
AC P58993;
                                                                                                                                                                                                                                                                                             Matches
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SEQUENCE FROM N.A.

STRATS_ATC 33993

KATNIS_ATC 33993

KATNIS_ATC 33993

KATNIS_ATC 33993

KATNIS_ATC 33993

KATNIS_ATC 3202145; PubMed=12024217; DOI=10.1038/4174599;

KA GA Silva A.C.R., Ferro J.A., Reinach F.C., Farah C.S., Furlan L.R.,

Alves L.M.C., do Amaral A.M., Bertolini M.C., Camargo L.B.A.,

Camarotte G., Cannavan F., Cardozo J., Chambergo F., Ciapina L.P.,

Raria J.B., Ferreira A.J.S., Ferreira R.C.C., Ferro M.I.T.,

Formighieri E.F., Franco M.C., Garegio C.C., Gruber A.,

Katsuyama A.M., Kishi L.T., Leite R.P., Lemos B.G.M., Lemos M.V.F.,

Martins E.C., Madianis J., Madeira A.M.B.N., Matyaki C.Y., Moon D.H.,

Moreira L.M., Novo M.T.M., Okura V.K., Oliveira M.C., Oliveira V.R.,

Moreira L.M., Rakia M.A., Tamura R.E., Teixeira B.C.,

Ropinola L.A.F., Takita M.A., Tanura R.S., Teixeira B.C.,

Spinola L.A.F., Takita M.A., Taulfi D., Tsai S.M., White F.F.,

Schubal J.C., Kitajima J.P.,

R. "Comparison of the genomes of two Xanthomonas pathogens with differing that. Appl. 441459-4432 20202 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN=306 / ATCC 13902 / XV 101;

SECREMBE ATCC 13902 / XV 101;

MEDLINE=22022145; PubMed=12024217; DOI=10.1038/417459a;

A da Silva A.C.R., Ferro J.A., Reinach F.C., Farah C.S., Furlan L.R.,

A Quaggio R.B., Monteiro-Vitorello C.B., Van Sluys M.A., Almeida N.F.,

A lves L.M.C., do Amaral A.M., Bertolini M.C., Camargo L.E.A.,

A cicarelli R.M.B., Coutinho L.L., Cursino-Santos J.R., El-borry H.,

Faria J.B., Ferreira A.J.S., Ferreira R.C.C., Ferro M.I.T.,

Katsuyama A.M., Kishi L.T., Leite R.P., Lemos E.G.M., Lemos M.V.F.,

Normighieri E.C., Machado M.A., Madelra A.M.B.N., Martinez-Rossi N.M.,

Martins E.C., Maidanis J., Menck C.F.M., Miyaki C.Y., Moon D.H.,

Moreira L.M., Novo M.T.M., Okura V.K., Oliveira M.C., Oliveira V.R.,

Pereira H.A., Rossi A., Sena J.A.D., Silva C., de Souza R.F.,

A Spinola L.A.F., Takita M.A., Tamura R.E., Teixeira E.C., Tezza R.I.D.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Xanthomonas axonopodis (pv. citri).
Bacteria, Proteobacteria, Gammaproteobacteria, Xanthomonadales,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 69; DB 2; Length 51; 10.8%; Pred. No. 5.1e+02; tive 33; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7A25C67893030E8F CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 CVVCGFLYHEADGIPEEGIAPGTRWEDVPDSWTCPDC 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            51 AA.
                                                                                                                                                                                                                                                                                                                                    GO; GO:0046872; F:metal ion binding; IEA.
GO; GO:0006118; P:electron transport; IEA.
InterPro; IPR004039; Rubredox:
InterPro; IPR001052; Rubredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                           PROSITE; PS00202; RUBREDOXIN, UNKNOWN_1.
PROSITE; PS50903; RUBREDOXIN_LIKE; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Name=rubA; OrderedLocusNames=XAC3414;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Xanthomonadaceae; Xanthomonas.
                                                                                                                                                                                                                                                                                                          EMBL; AE012443; AAM42538.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                Pfam; PF00301; Rubredoxin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Complete proteome.
SEQUENCE 51 AA; 5679 MW;
                                                                                                                                                                                                                                                                                                                                                                                                               ProDom; PD001610; Rubredox;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rubredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Q8PH46;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Q8PH46
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Trindade dos Santos M., Truffi D., Tsai S.M., White F.F.,
Setubal J.C., Kitajima J.P.;
"Comparison of the genomes of two Xanthomonas pathogens with differing
host specificities.";
host specificities.";
BMBL; AE011987; AAM38257.1;
EMBL; AE012887; IRO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Name=rub2; Synonyms=rbr-2; OrderedLocusNames=CT1101;
Chlorobium tepidum.
Bacteria; Chlorobi; Chlorobia; Chlorobiales; Chlorobiaceae;
Chlorobaculum.
                                                                                                                                                                                                                                                                                                                                                                                                                 DB 2; Length 51;
                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Pred. No. 5.1e+02;
33; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                     51 AA; 5706 MW; 6E26E27D93530E8F CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2 CVVCGFLYHEADGIPEEGIAPGTRWQDVPDTWTCPDC 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
28-CCT-2004 (Rel. 45, Last annotation update)
Rubredoxin 2 (Rd 2).
                                                                                                                                                             GO; GO:0046872; F:metal ion binding; IEA.
GO; GO:0006118; P:electron transport; IEA.
InterPro; IPR004039; Rubredox.
InterPro; IPR001052; Rubredoxin.
Pfam; PF00301; Rubredoxin: 1.
ProDom; PD001610; Rubredoxin: 1.
PROSITE; PS00202; RUBREDOXIN; UNKNOWN 1.
PROSITE; PS0903; RUBREDOXIN, LIKE; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  52 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 69;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                        10.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               4; Conservative
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Matches

RESULT 5

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21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
25-OCT-2004 (Rel. 45, Last annotation update)
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Iron.
Iron.
                                                                                                                                                                                                                                                                            MEDLINE=76231572; PubMed=938515;
                                                                                                                STANDARD;
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                                                                                                                                                                                   (Rd)
                                                                                                                                                                                                                                                                                              Bruschi M.;
                                                                                                             RUBR DESGI
P00270;
                                                                                                                                                                                  Rubredoxin
                                                                                                                                                                                                                                                                    SEQUENCE
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-!- FUNCTION: Rubredoxin is a small nonheme, iron protein lacking acid-labile sulfide. Its single Fe, chelated to 4 Cys, functions as an electron acceptor and may also stabilize the conformation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRAIN=ATCC 7956;
MEDLINE=91058526; PubMed=2244884;
Meyer J., Gagnon J., Sieker L.C., van Dorsselaer A., Moulis J.-M.;
Meyer John Clostridium thermosaccharolyticum. Amino acid
sequence, mass-spectrometric and preliminary crystallographic data.";
Biochem. J. 271:839-841 (1990).
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                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Direct protein sequencing; Electron transport; Iron; Metal-binding; Rubredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -1- COPACTOR: Binds 1 iron(III) ion per subunit (By similarity).
-1- SIMILARITY: Belongs to the rubredoxin family.
-1- SIMILARITY: Contains 1 rubredoxin-like domain.
PIR; A33173; A33173.
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              Prodom; PD001610; Rubredox; 1.
PROSITE; PS00202; RUBREDOXIN; 1.
PROSITE; PS50903; RUBREDOXIN LIKE; 1.
COMDISTE proteome; Electron Transport; Iron; Metal-binding; Bubredoxin. 1 57
                                                                                                                                                                                                             Length 52;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bacteria; Firmicutes; Clostridia; Thermoanaerobacteriales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Score 69; DB 1; Length 52; Similarity 10.8%; Pred. No. 5.2e+02; 4; Conservative 33; Mismatches 0; Indels
                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Clostridium thermosaccharolyticum (Thermoanaerobacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Iron (By similarity).
Iron (By similarity).
Iron (By similarity).
Iron (By similarity).
4FA4AlB0714E080F CRC64;
                                                                                                                                                                   Iron (By similarity).
4239E1BF9C52B203 CRC64
                                                                                                                 Iron (By similarity).
Iron (By similarity).
Iron (By similarity).
Iron (By similarity).
                                                                                                                                                                                                                                                                                 Score 69; DB 1;
Pred. No. 5.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Thermoanaerobacteriaceae; Thermoanaerobacterium
                                                                                                                                                                                                                                                                                                                                                                                           01-FEB-1991 (Rel. 17, Created)
01-FRB-1991 (Rel. 17, Last sequence update)
05-UUL-2004 (Rel. 44, Last annotation update)
Rubredoxin (Rd).
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                                                                                                                                                                                                      100.0%; Score 69; un
10.8%; Pred. No. 5.2e4
tive 33; Mismatches
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PROSITE; PS00202; RUBREDOXIN; 1.
PROSITE; PS50903; RUBREDOXIN_ILKE; 1.
InterPro; IPR001052; Rubredoxin.
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InterPro; IPR004039; Rubredox.
InterPro; IPR001052; Rubredoxin.
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PRINTS; PR00163; RUBREDOXIN.
             Pfam; PF00301; Rubredoxin; 1
PRINTS; PR00163; RUBREDOXIN.
                                                                                                                                                                                5746 MW;
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                                                                                                                                                                                52 AA;
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                                                                                                                                                                                                                                                                                                                                               RUBR_CLOTS
ID _RUBR_CLOTS
                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                           Query Match
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DOMAIN

Matches

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"Rubredoxin from Desulfovibrio gigas. A molecular model of the oxidized form at 1.4-A resolution.";
J. Mol. Biol. 197:525-541(1987).
-!- FUNCTION: Rubredoxin is a small nonheme, iron protein lacking acid-labile sulfide. Its single Fe, chelated to 4 Cys, functions as an electron acceptor and may also stabilize the conformation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -!- FUNCTION: Electron acceptor for cytoplasmic lactate dehydrogenase.
-!- COFACTOR: Binds 1 iron(III) ion per subunit.
-!- SUBCELLULAR LOCATION: Cytoplasmic.
-!- SIMILARITY: Belongs to the rubredoxin family.
-!- SIMILARITY: Contains 1 rubredoxin-like domain.
PIR; A00275; RUDVEG.
PDB; 1RBJ; NMX; A=1-52.
InterPro; IPR004039; Rubredox.
InterPro; IPR004039; Rubredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Desulfovibrio gigas.
Bacteria; Proteobacteria; Deltaproteobacteria; Desulfovibrionales;
Desulfovibrionaceae; Desulfovibrio.
NCBI_TaxID=879;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "The amino acid sequence of rubredoxin from the sulfate reducing bacterium, Desulfovibrio gigas."; Biochem. Biophys. Res. Commun. 70:615-621(1976).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           X-RAY CRYSTALLOGRAPHY (1.4 ANGSTROMS).
MEDLINE=88155649; PubMed=3441010;
Frey M., Sieker L.C., Payan F., Haser R., Bruschi M., Pepe G.,
le Gall J.;
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PRINTS; PR00163; RUBREDOXIN.
PRODOM; PD001610; RUBredox; 1.
PROSITE; P$00202; RUBREDOXIN; 1.
PROSITE; P$50903; RUBREDOXIN; 1.
PROSITE; P$50903; RUBREDOXIN; 1.
PROSITE; P$50903; RUBREDOXIN; 1.
PROSITE; P$50903; RUBREDOXIN; 1.
Notormylation; Iron; Metal-binding; Rubredoxin.
NOD RES 1 1 N-formylmethionine.
DOMĀIN 1 52 RUBredoxin-like.
37
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METAL
CONFLICT
STRAND
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PubMed=15077118; DOI=10.1038/nbt959; Heidelberg J.F., Seshadri R., Haveman S.A., Hemme C.L., Paulsen I.T., Kolonay J.F., Elsen J.A., Ward N.L., Methe B.A., Brinkac L.M., Daugherty S.C., DeBoy R.T., Dodson R.J., Durkin A.S., Madupu R., Nelson W.C., Sullivan S.A., Fouts D.E., Haft D.H., Selengut J., Peterson J.D., Davidsen T.M., Zafar N., Zhou L., Radune D., Dimitrov G., Hance M., Tran K., Khouri H.M., Gill J., Utterback T.R., Feldblyum T.V., Wall J.D., Voordow G., Fraser C.M.; "The genome sequence of the anaerobic, sulfate-reducing bacterium Desulfovibrio vulgaris Hildenborough.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 X-RAY CRYSTALLOGRAPHY (2 ANGSTROMS).
MEDLINE=7730456; PubMed=881725;
Adman E.T., Sieker L.C., Jensen L.H., Bruschi M., le Gall J.;
"A gtructural model of rubredoxin from Desulfovibrio vulgaris at 2-A
                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Non-heme iron proteins. The amino acid sequence of rubredoxin from Desulfovibrio vulgaris.";
Biochim. Biophys. Acta 434:4-17(1976).
                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
MEDLINE-88329747; PubMed=2843442; DOI=10.1016/0378-1119(88)90010-8;
                                                                                                                                                                                                                                                                                                                                                                          rubredoxin (rub) and a putative rubredoxin oxidoreductase (rbo) in Desulfovibrio vulgaris Hildenborough."; J. Bacteriol. 171:4996-5004(1989).
                                                                                                                                                                                                                                                                                     Bacteria; Proteobacteria; Deltaproteobacteria; Desulfovibrionales; Desulfovibrionaceae; Desulfovibrio. NCBI_TaxID=882;
                                                                                                                                                                                                                                                     Name=rub; OrderedLocusNames=DVU3184;
Desulfovibrio vulgaris (strain Hildenborough / ATCC 29579 / NCIMB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=76232334; PubMed=7308; DOI=10.1016/0005-2795(76)90030-1;
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MEDLINE-89359139; PubMed=2549009;
Brumlik M.J., Vocrdouw G.; "Analysis of the transcriptional unit encoding the genes for "Analysis of the transcriptional unit
                                                           Length 52;
                                                                                Indels
                                   3E08147A4AC262F7 CRC64;
                                                                                                                    6 CTVCGYEYDPAKGDPDSGIKPGTKFEDLPDDWACPVC 42
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                                                        100.0%; Score 69; DB 1;
10.8%; Pred. No. 5.2e+02;
tive 33; Mismatches 0
                                                                                                                                                                                                        21-JUL-1986 (Rel. 01, Created)
01-FEB-1991 (Rel. 17, Last sequence update)
25-OCT-2004 (Rel. 45, Last annotation update)
                                                                                                                                                                                   52 AA
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MEDLINE=91124457; PubMed=1992166;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nat. Biotechnol. 22:554-559(2004)
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                                   5677 MW;
                                                                                Conservative
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                                                       Ouery Match
Best Local Similarity
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AA;
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P00269;
                                   SEQUENCE
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                                                                                                   X-YAY CRYSTALLOGRAPHY (0.92 ANGSTROMS).

Dauter Z., Butterworth S., Sieker L.C., Sheldrick G., Wilson K.S.;
Submitted (DEC-1997) to the PDB data bank.

-!- FUNCTION: Rubredoxin is a small nonheme, iron protein lacking acid-labile sulfide. Its single Fe, chelated to 4 Cys, functions as an electron acceptor and may also stabilize the conformation of
                                                                                                                                                                                                                                                   FUNCTION: Electron acceptor for cytoplasmic lactate dehydrogenase. COFACTOR: Binds 1 iron(III) ion per subunit. SUBCELLUIAR LOCATION: Cytoplasmic. SIMILARITY: Belongs to the rubredoxin family. SIMILARITY: Contains 1 rubredoxin-like domain.
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Adman E.T., Seiker L.C., Jensen L.H.; "Structure of rubredoxin from Desulfovibrio vulgaris at 1.5-A
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Short protein sequencing;

Electron transport; Iron; Metal-binding; Rubredoxin.

Rubredoxin-like.
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10.8%; Pred. No. 5.2e+02;
ive 33; Mismatches 0;
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PRINTS; PR00163; RUBREDOXIN.
ProDom; PD001610; Rubredoxi.
PROSITE; PS00202; RUBREDOXIN; 1.
PROSITE; PS50903; RUBREDOXIN_LIKE; 1.
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                                                           Mol. Biol. 217:337-352(1991).
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InterPro; IPR001052; Rubredox:
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EMBL; AE017320; AAS97654.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; AE017320; AAS97654.1;
PIR; B33962; RUDV.
PDB; 1RB9; X-ray; @=1-52.
PDB; 7RXN; X-ray; @=1-52.
PDB; 8RXN; X-ray; A=1-52.
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Best Local Similarity 10.00
-204 4; Conservative
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MEDLINE-99190993; PubMed=10089348; DOI=10.1107/S0907444998011810;

Misaki S., Morimoto Y., Ogata M., Yagi T., Higuchi Y., Yasuoka N.;

Misaki S., Morimoto Y., Ogata M., Yagi T., Higuchi Y., Yasuoka N.;

"Structure determination of rubredoxin from Desulfovibrio vulgaris

Miyazaki F in two crystal forms.";

Acta Crystallogr. DS:408413(1999).

-!- Crystallogr. DS:408413(1999).

-!- FUNCTION: Rubredoxin is a small nonheme, iron protein lacking

acid-labile sulfide. Its single Fe, chelated to 4 Cys, functions

as an electron acceptor and may also stabilize the conformation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           -!- FUNCTION: Electron acceptor for cytoplasmic lactate dehydrogenase.
                                                                                                                                                                                                                                                                                                                                                                        STRAIN=150Jate F; MEDLINE=97236449; PubMed=9116039; DOI=10.1016/S0167-4781(96)00203-5; Kitamura M., Koshino Y., Kamikawa Y., Kohno K., Kojima S., Miura K., Sagara T., Abutsu H., Kumagai I., Nakaya T.; "Cloning and expression of the rubredoxin gene from Desulfovibrio vulgaris (Miyazaki F) -- comparison of the primary structure of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=90234754; PubMed=2561345; DOI=10.1016/0300-9084(89)90020-5; Shimuzu F., Ogata M., Yagi T., Wakabayashi S., Matsubara H.; "Amino acid sequence and function of rubredoxin from Desulfovibrio
                                                                                                                                                                                                                                                                            Bacteria; Proteobacteria; Deltaproteobacteria; Desulfovibrionales;
Desulfovibrionaceae; Desulfovibrio.
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PROSITE; PS50903; RUBREDOXIN_LIKE; 1.
3D-structure; Direct protein_sequencing; Electron transport;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          -!- SIMILARITY: Belongs to the rubredoxin family.
-!- SIMILARITY: Contains 1 rubredoxin-like domain.
01-APR-1990 (Rel. 14, Created)
01-APR-1990 (Rel. 14, Last sequence update)
25-OCT-2004 (Rel. 45, Last annotation update)
                                                                                                                  52 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Biochim. Biophys. Acta 1351:239-247(1997).
                                                                                                                                                                                                                                                         Desulfovibrio vulgaris (strain Miyazaki)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -!- SUBCELLULAR LOCATION: Cytoplasmic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PDB; IRDV; X-ray; @=1-52.
PDB; ZRDV; X-ray; A/B/C=1-52.
InterPro; IPR004039; Rubredox.
InterPro; IPR001052; Rubredoxin.
Pfam; PF00301; Rubredoxin; 1.
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ProDom; PD001610; Rubredox; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Biochimie 71:1171-1177(1989).
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                                                                                                                    STANDARD;
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                                                                                                                                                                        01-APR-1990 (Rel
25-OCT-2004 (Rel
Rubredoxin (Rd).
                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=883;
                                                                                                                    RUBR DESVM
                                                                                                                                                                                                                                         Name=rub;
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SEQUENCE, AND MASS SPECTROMETRY.
MEDLINE=95243660; PubMed=7726577; DOI=10.1006/abbi.1995.1207;
Lee W.Y., Brune D.C., Lobrutto R., Blankenship R.E.;
"Isolation, characterization, and primary structure of rubredoxin from the photosynthetic bacterium, Heliobacillus mobilis.";
Arch. Blochem. Biophys. 318:80-88(1995).
-!- FUNCTION: Rubredoxin is a small nonheme, iron protein lacking acid-labile sulfide. Its single Fe, chelated to 4 Cys, functions as an electron acceptor and may also stabilize the conformation of
                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteria; Firmicutes; Clostridia; Clostridiales; Heliobacteriaceae;
Heliobacillus.
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ProDom; PD001610; Rubredox; 1.
PROSITE; PS00202; RUBREDOXIN; 1.
PROSITE; PS50903; RUBREDOXIN LIKE; 1.
Direct protein sequencing; Electron transport; Formylation; Iron; Metal-binding; Rubredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the molecule.
-!- COFACTOR: Binds 1 iron(III) ion per subunit (By similarity).
-!- MASS SPECTROMETRY: MW=5671.3; METHOD=MALDI; RANGE=1-52;
                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                      Length 52;
                                                                                                                                                                                                                                                                                  %; Score 69; DB 1; Length 52
; Pred. No. 5.2e+02;
33; Mismatches 0; Indels
                                                                                                                                                                                                                                                               6443741A8A8063A2 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -!- SIMILARITY: Belongs to the rubredoxin family.
-!- SIMILARITY: Contains 1 rubredoxin-like domain.
PIR; 855620; 865620.
HSSP; P15412; ZRDV.
                                                                                                                                                                                                                                                                                                                                               |::|::::::::|::|
CTVCGYEYDPAEGDFDNGVKPGTAFEDVPADWVCPIC 42
                                                                                                                                                                                                                                                                                                                                  N-formylmethionine.
Rubredoxin-like.
Metal-binding; Rubredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                    15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                               PRT;
                               Iron.
                                            Iron.
Iron.
Iron.
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InterPro; IPR001052; Rubredoxin.
Pfam; PF00301; Rubredoxin; 1.
                                                                                                                                                                                                                                                                                   100.08;
                                                                                                                                                                                                                                                               5598 MW;
                                                                                                                                                                                                                                                                                               10.8%;
                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 10.00
Fra 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                 STANDARD;
                     Heliobacillus mobilis.
Formylation; Iron;
                                                                                                                                                                                                                                                              52 AA;
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NCBI_TaxID=1521;
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                                                                                                                   SEQUENCE
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Q9X709
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN=Nichols;
MEDLINE=98332770; PubMed=9665876; DOI=10.1126/science.281.5375.375;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Treponema pallidum.
Bacteria; Spirochaetes; Spirochaetales; Spirochaetaceae; Treponema
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PROSITE; PS50903; RUBREDOXIN LIKE; 1.
Complete proteome; Electron Transport; Iron; Metal-binding;
Rubredoxin. 1 52 Rubredoxin-like.
                                                                                                                                                                                                                                   %; Score 69; DB 1; Length 52;
; Pred. No. 5.2e+02;
33; Mismatches 0; Indels
                                                                                                                                                  Lron (By similarity):
659241131D3124AD CRC64;
N-formylmethionine.
Rubredoxin-like.
Iron (By similarity).
Iron (By similarity).
                                                                                                                            similarity).
                                                                                                                                                                                                                                                                                                                                                                                                  6 CLVCGYVYDDAKGDPDHGIAPGTAFEDLPADWVCPLC 42
                                                                                                                                                                                                                                                                                                                                                             (Rel. 37, Last sequence update) (Rel. 45, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           52 AA
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InterPro; IPR001052; Rubredoxin.
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PRINTS; PR00163; RUBREDOXIN.
                                                                                                                                                                                                                                             100.08;
                                                                                                                                                                              5619 MW;
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Best Local Similarity lu...
A; Conservative
A; Conservative
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25-OCT-2004
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   MOD_RES
DOMAIN
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EMBL; AVE49956; AAS55565.1; -

R GWG GO:0005576; C:extracellular; IEA.

GO; GO:000552; P:defense response; IEA.

GO; GO:000562; P:response to pest, pathogen or parasite; IEA.

INTERPO: IPRO06081; Defensin_alpha.

InterPo: IPR006081; Defensin_alpha.

InterPo: IPR006081; Defensin_alpha.

INTERPO: IPR001007; VWF C.

INTERPO: IPR001007;
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Bacteria; Firmicutes; Clostridia; Clostridiales; Clostridiaceae;
Clostridium.
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Drosophila mauritiana (Fruit fly).
Bukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                            DB 1; Length 52;
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to the EMBL/GenBank/DDBJ databases.
Iron (By similarity).
Iron (By similarity).
Iron (By similarity).
Iron (By similarity).
A76C9F012A70FIEI CRC64;
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Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-JUL-2004 (TrEMBLrel. 27, Last sequence update) 05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                          6 CDLCGWGYNPEVGDADGGIPAGTAFENLPDHWECPLC 42
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10.8%; Pred. No. 5.2e+02;
                                                                                                                                                                                                                                                                   Pred. No. 5.2e+02;
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                                                                                                                                                                                                                                                                                                      33; Mismatches
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                                                                                                                                                                                                                                Score 69;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Created)
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5568 MW;
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                                                                                                                                                                                                                                                                       10.8%;
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Best Local Similarity
Matches 4; Conserva
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Best Local Similarity
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52 AA;
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STRAIN=ATCC 35405 / DSM 14222;

RubMed=15064399; DOI=10.1073/pnas.0307639101;

R Seshadri R., Myers G.S.A., Tettelin H., Elsen J.A., Heidelberg J.F.,

B Seshadri R., Myers G.S.A., Tettelin H., Elsen J.A., Heidelberg J.F.,

B Davidsen T.M., Dabloy R.T., Fours D.E., Haft D.H.,

Selengut J., Ren Q., Brinkac L.M., Madupu R., Kolonay J.F.,

B Durkin S.A., Daugherty S.C., Shetty J., Shvartsbeyn A.,

R Gebregorgis E., Geer K., Tsegaye G., Malek J.A., Ayodeji B.,

R Shatsman S., McLeod M.P., Smajs D., Howell J.K., Pal S., Amin A.,

A Shatsman S., McLeod M.P., Smajs D., Howell J.K., Pal S., Amin A.,

R Wahisth P., Moris S.J., Fraser C.M., Paulsen I.T.,

T "Comparison of the genome of the oral pathogen Treponema denticola

With other spirochete genomes.",

B Proc. Natl. Acad Sci. US.A. 101.5646-5651(2004).
Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Treponema denticola.
Bacteria; Spirochaetes; Spirochaetales; Spirochaetaceae; Treponema.
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05-JUL_2004 (TrEMBLrel. 27, Last sequence update)
05-JUL_2004 (TrEMBLrel. 27, Last annotation update)
05-JUL_2004 (TrEMBLrel. 27, Last annotation update)
Rubredoxin.
05-GeredLocusNames=GSU3188;
Geobacter sulfurreducens.
Bacteria; Proteobacteria; Deltaproteobacteria; Desulfuromonadales;
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ProDom; PD001610; Rubredox; 1.
PROSITE; PS00202; RUBREDOXIN; 1.
PROSITE; PS50903; RUBREDOXIN_LIKE; 1.
Complete proteome; Electron transport; Iron; Metal-binding; Rubredoxin.
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   0; Indels
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                                                                                                                                                                                                                                                                                                                                                                      Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Proc. Natl. Acad. Sci. U.S.A. 101:5646-5651(2004).
-!- SIMILARITY: Belongs to the rubredoxin family.
-!- SIMILARITY: Contains 1 rubredoxin-like domain.
EMBL; AE017249; AAS11541.1; -.
HSSP; P00268; 1B13.
TIGR; TDE1052; -.
                                                                                                CDLCGYYYDPALGDPDGGIAPGTAFEDIPDDWVCPLC 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     52 AA.
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33; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        33; Mismatches
                                                                                                                                                                                                                                                                                                                                            Created)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        InterPro; IPR004039; Rubredox.
InterPro; IPR001052; Rubredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pfam; PF00301; Rubredoxin; 1.
                                                                                                                                                                                                                                                                                                                                                               05-JUL-2004 (TrEMBLrel, 27, 05-JUL-2004 (TrEMBLrel, 27,
                                                                                                                                                                                                                                                                                                                                         (TrEMBLrel. 27,
                                                                                                                                                                                                                                                                                                                                                                                                                                                             OrderedLocusNames=TDE1052;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4; Conservative
4; Conservative
                                                                                                                                                                                                                                                                               PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=158;
                                                                                                                                                                                                                                                                                                                                            05-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                   Rubredoxin.
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Q747S7;
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                                                                                                                                                                                                              RESULT 14
Q73NV0
Matches
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MEDLINE=21992816; PubMed=11997336; DOI=10.1101/gr.219302;
Bao Q., Tian X., Li W., Xu Z., Xuan Z., Hu S., Dong W., Yang J.,
Chen Y., Xue Y., Xu Y., Lai X., Huang L., Dong X., Ma Y., Ling L.,
Tan H., Chen R., Wang J., Yu J., Yang H.;
"A complete sequence of the T. tengcongensis genome.";
Genome Res. 12:689-700(2002).
-!- SIMILARITY: Belongs to the rubredoxin family.
-!- SIMILARITY: Contains 1 rubredoxin-like domain.
EMBL; AR031363; AAM25316.1; --
HSSP; P00270; IRDG.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
100.0%; Score 69; DB 2; Length 52;
Best Local Similarity 10.8%; Pred. No. 5.2e+02;
Matches 4; Conservative 33; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Thermoanaerobacter tengcongensis.
Bacteria; Firmicutes; Clostridia; Thermoanaerobacteriales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Score 69; DB 2; Length 52; 10.8%; Pred. No. 5.2e+02;
                                                                                                          Submitted (FEB-2001) to the EMBL/GenBank/DDBJ databases.
-!- SIMILARITY: Belongs to the rubredoxin family.
-!- SIMILARITY: Contains 1 rubredoxin-like domain.
EMBL; AJ338458; CAB41597.1; -.
                                                                                                                                                                                                                                                               GO; GO:0005489; F:electron transporter activity; IEA. GO; GO:0046872; F:metal ion binding; IEA. GO; GO:0006118; P:electron transport; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PROSITE; PS00202; RUBREDOXIN; 1.
PROSITE; PS50903; RUBREDOXIN LIKE; 1.
Electron transport; Iron; Metal-binding; Rubredoxin.
SEQUENCE 52 AA; 5658 MW; AF0F517D000E2FED CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | ::| :::::::| | CTVCGYVYDPEEGDPDGGIAPGTAFEDIPEDWVCPLC 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Last sequence update)
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Thermoanaerobacteriaceae; Thermoanaerobacter.
NCBI_TaxID=119072;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               52 AA.
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PROSITE; PS50903; RUBREDOXIN LIKE; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-JUN-2002 (TrEMBLrel. 21, Created)
01-JUN-2002 (TrEMBLrel. 21, Last seq
01-OCT-2003 (TrEMBLrel. 25, Last ann
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                                                                                                                                                                                                                                                                                                                                                            InterPro; IPR004039; Rubredox.
InterPro; IPR001052; Rubredoxin.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRINTS; PR00163; RUBREDOXIN.
ProDom; PD001610; Rubredox; 1.
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PRINTS; PR00163; RUBREDOXIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                Pfam; PF00301; Rubredoxin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OrderedLocusNames=TTE2154;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRELIMINARY;
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                      SEQUENCE FROM N.A. STRAIN=ATCC 35319;
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us-09-763-616-2.dx.rup

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0;
                                                 STRAIN=PCA / ATCC 51573;

STRAIN=PCA / ATCC 51573;

PtubMed-14671304; DOI=10.1126/science.1088727;

Methe B.A., Nelson K.E., Eisen J.A., Paulsen I.T., Nelson W.C., Heidelberg J.F., Wu D., Wu M., Ward N.L., Beanan M.J., Dodson R.J., Madupu R., Brinkac L.M., Daugherty S.C., DeBoy R.T., Durkin A.S., Gwinn M.L., Kolonay J.F., Sullivan S.A., Haft D.H., Selengut J., Davidsen T.M., Zafar N., White O., Tran B., Romero C., Forberger H.A., Weidman J.F., Khouri H.M., Feldblyum T.V., Utterback T.R., Wan Aken S.E., Lovley D.R., Fraser C.M.;
"Genome of Geobacter sulfurreducens: metal reduction in subsurface
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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PROSITE; PS00202; RUBREDOXIN; 1.
PROSITE; PS50903; RUBREDOXIN_LIKE; 1.
Complete proteome; Blectron transport; Iron; Metal-binding; Rubredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 69; DB 2; Length 52; 10.8%; Pred. No. 5.2e+02; ive 33; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                   GO; GO:0005489; F:electron transporter activity; IEA. GO; GO:0046872; F:metal ion binding; IEA. GO; GO:0006118; P:electron transport; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   52 AA; 5877 MW; 5480469CA72044F6 CRC64;
                                                                                                                                                                                                                                                                                   Science 302:1967-1969(2003).
-!- SIMILARITY: Belongs to the rubredoxin family.
-!- SIMILARITY: Contains 1 rubredoxin-like domain.
EMBL, AE017180; ARR36579.1; -.
HSSP: P24297; IUUS.
TIGR; GSU3188; -.
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InterPro; IPR001052; Rubredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pfam; PF00301; Rubredoxin; 1
PRINTS; PR00163; RUBREDOXIN.
Geobacteraceae; Geobacter.
NCBI_TaxID=35554;
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Best Local Similarity 10.8*
Matches 4; Conservative
                                                                                                                                                                                                                                                                   environments.";
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Search completed: May 27, 2005, 00:35:22 Job time : 176 secs

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Copyright (c) 1993 - 2005 Compugen Ltd.
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Maximum Match 100%
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Gapop 10.0 , Gapext 0.5
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69
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Result No.	Score	Query Match	Length	DB	OI.	Description
1	69		37	1	PCT-US99-17130-287	287,
7 Y	69	100.0	37	220	US-09-763-616-2	Sequence 2, Appli
J 47	9 9		38	U [-	US-10-331-334-234 US-60-273-808-33	33,
2	69		38	_	US-60-278-983-33	33,
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15	69		38		US-08-467-459-40	40,
16	69	-	39		US-08-467-459A-40	40,
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16	69		9 6		US-08-467-906-40	40,
20	69		39		US-08-468-731-40	40,
21	69	-	39		US-08-468-731A-40	40,
22	69 6	_	33		US-08-469-526-40	40
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25	69		9 6		US-08-470-335-40	40,
26	69	_	39		US-08-470-335B-40	40,
27	69		66		US-08-471-148-40	40,
8 6	5 G		20.0		US-08-471-833-40	4 4
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31	69		36		US-08-472-008-40	4
32	69		39		US-08-472-065-40	40,
33	69		66		US-08-472-065F-40	40,
ω . 4. r	5 6		9 0		US-08-535-200-40	0,4
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) (A)	69		3 6	1 7	US-08-734-665-40	40,
38	69		33	_	US-08-734-665A-40	40,
39	69		39	~	US-08-734-665-40	40,
40	69		9 9	Н,	US-08-734-666-40	40,
4.4	א ס		א מ		118-08-736-010-40	, 0
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44	69		39	~	US-08-736-070A-40	40,
45	69		39	а .	US-08-736-070-40	40,
					CHARTMAN + va	
					ALLGINEENIO	
RESULT 1						
\sim	-17130-	17130-287	Ş	00011111/00	0 6 17 100 01	
; GENERAL 1	L INFOR	APPILC MATION:	3	7	CT / T	
	Ε.	G G	поше	cienc	es, Inc.	
	TITLE OF INVENT	ENTION:	98 Human	lan S	ecre	
, CURRE	NT APPL	TION	; ₽		PCT/US99/17130	
	NT FILE	NG DATE	1999-	0.4	,	
	ER APPL	ICATION NO DATE	NUMBER	200	1/094,657	
	ER APPL	ICATION	NUMBER: 6		0/095,486	
, EARLI	ER FILI	NG DATE	1998-			
, EARLI	ER PILI	NG DATE	1998-		0/095,455 06	
; EARLIER	ER APPL	APPLICATION	NUMBER: 6		0,095,454	
; EARLI	ER FILI	NG DATE	1998-	-80	,	
FAKLI	ER APPL	ICATION	NUMBER	9 6	60/096,319	

; EARLIER FILING DATE: 1998-08-12

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INVENTION: 98 Human Secreted Proteins
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Description of Artificial Sequence: Illustrative OTHER INFORMATION: zinc domain peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
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                                                                                                                                                           Query Match 100.0%; Score 69; DB 1; Length 37; Best Local Similarity 10.8%; Pred. No. 3.2e+03; Matches 4; Conservative 33; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: TAN, YEE JOO
APPLICANT: BERRHEIDE, WALTER
APPLICANT: BERRHEIDE, WALTER
APPLICANT: SIM, ANTHONY EUGENE
APPLICANT: SIM, MUI MUI
TITLE OF INVENTION: THERAPEUTIC COMPOUNDS AND METHODS
FILE REPREBRICE: BERN3001/JDB
CURRENT APPLICATION NUMBER: US/99/763,616
CURRENT APPLICATION NUMBER: PCT/AU99/00724
PRIOR FILING DATE: 1999-09-03
PRIOR FILING DATE: 1999-00-03
PRIOR FILING DATE: 1999-00-15
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100.0%; Pred. No. 3.2e+03;
tive 0; Mismatches 0;
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LOCATION: (35)...(36)
OTHER INFORMATION: Variable amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: MOD RES
LOCATION: (5)...(33)
OTHER INFORMATION: Variable amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: (2)...(3)
OTHER INFORMATION: Variable amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: AU PP5733/98 PRIOR FILING DATE: 1998-09-04 NUMBER OF SEQ ID NOS: 2 SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                     Sequence 2, Application US/09763616 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: BERNARD, HANS-ULRICH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
                    PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.
Matches 37; Conservative
    NUMBER OF SEQ ID NOS: 364
                                                                                                  ORGANISM: Homo sapiens
PCT-US99-17130-287
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: MOD_RES
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US-09-763-616-2
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US-09-763-616-2
                       SOFTWARE: Pa
                                                                                   TYPE: PRT
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Sequence 294, Application US/10351334; GENERAL INFORMATION:
, APPLICANT: Komatsoulis et al

RESULT 3 US-10-351-334-294

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Sequence 33, Application US/60278983
Sequence 33, Application US/60278983
SEQUENCE INFORMATION:
APPLICANT: Bristol-Myers Squibb Company
TITLE OF INVENTION: POLYNUCLEOTIDE ENCODING A NOVEL HUMAN G-PROTEIN COUPLED RECEPTOR:
TITLE OF INVENTION: HOPREMY27, EXPRESSED HIGHLY IN THE SPLEEN AND TESTIS
FILE REFERENCE: D0134.pspl.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      G-PROTEIN COUPLED RECEPTOR
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TITLE OF INVENTION: POLYNUCLEOTIDE ENCODING A NOVEL HUMAN G-PROTEIN COUPLE
TITLE OF INVENTION: HOFRBMY27, EXPRESSED HIGHLY IN THE SPLEEN AND TESTIS
FILE REFERENCE: D0134
CURRENT FILING DATE: 2001-03-07
NUMBER OF SEQ ID NOS: 34
SOFTWARE: Patentin version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 37;
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; Pred. No. 3.2e+03;
33; Mismatches 0;
CURRENT APPLICATION NUMBER: US/10/351,334
CURRENT APPLICATION NUMBER: US/10/351,334
CURRENT FILING DATE: 2003-01-27
PRIOR APPLICATION NUMBER: 09/489,847
PRIOR PLING DATE: 2002-01-25
PRIOR APPLICATION NUMBER: 09/489,847
PRIOR PLING DATE: 2000-01-24
PRIOR PLING DATE: 1999-07-29
PRIOR PLING DATE: 1999-07-29
PRIOR PLING DATE: 1998-06-25
PRIOR APPLICATION NUMBER: 60/095,486
PRIOR APPLICATION NUMBER: 60/095,486
PRIOR PLING DATE: 1998-08-05
PRIOR PLING DATE: 1998-08-15
PRIOR FILING DATE: 1998-08-15
PRIOR FILING DATE: 1998-08-15
PRIOR PLILNG DATE: 1998-08-16
PRIOR PLILNG DATE: 1998-08-16
PRIOR APPLICATION NUMBER: 60/095,455
PRIOR PLILNG DATE: 1998-08-06
PRIOR APPLICATION NUMBER: 60/095,455
PRIOR PLILNG DATE: 1998-08-06
PRIOR PLILNG DATE: 1998-08-06
PRIOR APPLICATION NUMBER: 60/095,455
PRIOR PLILNG DATE: 1998-08-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 33, Application US/60273808; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 10.8%;
Matches 4; Conservative 33.
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US-60-273-808-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; ORGANISM: Homo sapiens
US-10-351-334-294
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LENGTH: 38
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100.0%; Score 69; DB 4; Length 39; 10.8%; Pred. No. 3.4e+03; ive 33; Mismatches 0; Indels
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MEDIUM TYPE: Diskette, 5.25 inch, 360 kb storage
COMPUTER: IBM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3 CGCCKCCRTTCACRCAGAAGGTCTTCTCCTTCTCAGC 39
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/984,085
FILING DATE: 01-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/951,747
FILING DATE: 25-SEP-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/927,337
ATTORNEY AGENT INFORMATION:
NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 04585/017004
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 40, Application US/08209204A
GENERAL INFORMATION:
APPLICANT: Robert Sklar, Mark Marchionni,
APPLICANT: David I. Gwynne
TITLE OF INVENTION: METHODS FOR TREATING
NUMBER OF SEQUENCES: 184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04585/028001
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/209,204A
FILING DATE: 08-MAR-1994
CLASSIFICATION: 514
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APPLICATION NUMBER:
APPLICATION NUMBER:
06-MAY-93
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
      29-JAN-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 10.55
Thes 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: Massachusetts
ZIP: 02110-2804
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-011-396A-40
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US-08-209-204A-40
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GENERAL INFORMATION:
APPLICANT: Kumar, Manoj
APPLICANT: Mazeaud, Isabelle
APPLICANT: Christiano, Steven P.
TITLE OF INVENTION: Controlled Release of Active Agents Utilizing Repeat Sequence
TITLE OF INVENTION: Protein Polymers
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Pred. No. 3.3e+03;
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Sequence 40, Application US/08011396A

GENERAL INFORMATION:

APPLICANT: GENERAL INFORMATION:

TILLE OF INVENTION: INHIBITORS OF CELL PROLIFERATION,

TILLE OF INVENTION:

STREET: 225 Franklin Street

CITY: Boston

STATE: Massachusetts

INFO 1221-2804

COMPUTER: Diskette, 5.25 inch, 360 kb storage
                                                                                                                                                                                                                                          Indels
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CURRENT APPLICATION NUMBER: PCT/US04/15318
CURRENT APPLICATION NUMBER: US 60/470,465
PRIOR APPLICATION NUMBER: US 60/470,465
PRIOR FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 31
SOFTWARE: Patentin version 3.2
SEQ ID NO 22
LENGTH: 39
                                                                                                                                                                                                                                                                                  2 CAGCAGCGCCGCATGTACAACGGGTCGTGCTGCCGC 38
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                                                                                                                                                                                                                                       33; Mismatches
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    2001-03-27
                                                                                                                                                                                           Query Match
Best Local Similarity 10.8%; Ematches 4; Conservative 33;
CURRENT FILING DATE: 2001-03-2
NUMBER OF SEQ ID NOS: 34
SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: PC-DOS
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Matches 4; Conservative
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OTHER INFORMATION: rspp
                                                         SEQ ID NO 33
LENGTH: 38
TYPE: PRT
ORGANISM: homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: unknown
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PCT-US04-15318-22
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                                                                                                                                                   US-60-278-983-33
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APPLICANT: Sklar, Robert
APPLICANT: Marchionni, Mark
APPLICANT: Marchionni, Mark
APPLICANT: Gwynne, David
TITLE OF INVENTION: Misorders
TITLE OF INVENTION: Disorders
FILE REPRENCE: 04585/028001
CURRENT APPLICATION NUMBER: US/08/209,204E
CURRENT FILING DATE: 1994-03-08
PRIOR APPLICATION NUMBER: US 08/059,022
PRIOR PELLING DATE: 1993-05-06
NUMBER OF SEQ ID NOS: 420
SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 40
LENGTH: 39
                   APPLICANT: Marchionni, Mark
APPLICANT: Gwynne, David
TITLE OF INVENTION: Methods for Treating Muscle Diseases and
TITLE OF INVENTION: Disorders
FILE REFERENCE: 04585/028001
CURRENT APPLICATION NUMBER: US/08/209,204D
CURRENT FILING DATE: 1994-03-18
PRIOR FILING DATE: 1993-05-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 39;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       |::|::::::::|::|cgcckccrtrckcagaaggrctrcrccrtrcrcagc 39
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10.8%; Pred. No. 3.4e+03;
tive 33; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 69; DB 6;
10.8%; Pred. No. 3.4e+03;
iive 33; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Robert Sklar, Mark Marchionni,
APPLICANT: David I. Gwynne
TITLE OF INVENTION: METHODS FOR TREATING
NUMBER OF SEQUENCES: 184
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 417
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-209-204E-40; Sequence 40, Application US/08209204E; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 12
US-08-461-097-40
; Sequence 40, Application US/08461097
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Lag 4; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Bos taurus
US-08-209-204E-40
                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT ORGANISM: Bos taurus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-209-204D-40
                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                Length 39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                 0; Indels
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                                                                                                                                                                                                                                                                                                               APPLICANT: Sklar, Robert
APPLICANT: Marchionni, Mark
APPLICANT: Gwynne, David
TITLE OF INVENTION: METHODS FOR TREATING MUSCLE
TITLE OF INVENTION: DISEASES AND DISORDERS
NUMBER OF SEQUENCES: 379
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 69; DB 6;
Pred. No. 3.4e+03;
3; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READBLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OCREATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/209,204B
FILING DATE: 08-MAR-1994
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/059,022
                                                                                                                                                                                100.0%; Score 69; DB 6;
10.8%; Pred. No. 3.4e+03;
trive 33; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04585/028001
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FILING DATE: 06-MAX-1933
ATTORNEY/AGENT INFORMATION:
NAME: Bleker-Brady, Kristina
REGISTRATION NUMBER: 39,109
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-428-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 40, Application US/08209204B GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Clark & Elbing LLP
STREET: 176 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      33;
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Best Local Similarity 10.8%;
Matches 4; Conservative 33,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 39 amino acids
                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           617-428-7045
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                          TYPE: amino acid
STRANDEDNESS:
TOPOLOGY: linear
US-08-209-204A-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amino acid
                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Boston
STATE: MA
COUNTRY: USA
LENGTH: 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02110
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TELEX:
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Gaps . 0 ö

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CITY: Boston STATE: Massachusetts ZIP: 02110-2804 COMPUTER READABLE FORM:

US-08-209-204D-40; Sequence 40, Application US/08209204D; GENERAL INFORMATION:

Gaps

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GENERAL INFORMATION:
APPLICANT: GWynne, David I.; Marchionni, Mark;
APPLICANT: McBurney, Robert N.
TITLE OF INVENTION: THEIR PREPARATION AND USE
NUMBER OF SEQUENCES: 184
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Boston
STATE: Massachusetts
ZIP: 0211-2804
COMPUTER RRADABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 360 kb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3 CGCCKCCRITCACRCAGAAGGTCTTCTCCTTCTCAGC 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04585/028002
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: PC-DOS
SOFTWARE: Wordperfect
STORENT APPLICATION DATA:
APPLICATION NUMBER: US/08/466,446
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PILING DATE:
CLASSIFICATION: 436
PILING DATE:
APPLICATION DATA:
APPLICATION DATA:
FILING DATE: 29-JAN-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/984,085
PILING DATE: 01-DEC-1992
PRIOR APPLICATION NUMBER: 07/951,747
PILING DATE: 25-SEP-1992
PRIOR APPLICATION NUMBER: 07/951,747
PILING DATE: 10-ACC-1992
PRIOR APPLICATION NUMBER: 07/951,747
PILING DATE: 10-ACC-1992
APPLICATION NUMBER: 07/951,747
PILING DATE: 10-ACC-1992
ALTORNEY/AGENT INFORMATION:
      REFERENCE/DOCKET NUMBER: 04
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-428-0200
TELEPAX: 617-428-7045
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
                                                                                                                                        TELEX:
; INFORMATION FOR SEQ ID NO: 4
; SEQUENCE CHARACTERISTICS:
; LENGTH: 39 amino acids
; TYPE: amino acid STRANDEDNESS: single
; TOPOLLGY: linear
US-08-461-097A-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity 10.89
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM
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US-08-466-446-40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Sklar, Robert
APPLICANT: Marchionni, Mark
APPLICANT: Gwynne, David
TITLE OF INVENTION: METHODS FOR TREATING MUSCLE
TITLE OF INVENTION: DISEASES AND DISORDERS
NUMBER OF SEQUENCES: 383
CORRESPONDENCE ADDRESS:
ADDRESSEE: Clark & Elbing LLP
STREET: 176 Federal Street
CITY: Boston
Diskette, 5.25 inch, 360 kb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3 CGCCKCCRTTCACRCAGAAGGICTTCTCCTTCTCAGG 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match

100.0%; Score 69; DB 8;
Best Local Similarity 10.8%; Pred. No. 3.4e+03;
Matches 4; Conservative 33; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/08/461,097A
FILING DATE: 05-JUN-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/059,022
FILING DATE: 06-MAY-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/209,204
FILING DATE: 08-MAR-1994
ATTORNEY,AGENT THEORMATION:
AMDER: APPLICATION:
AMDER: APPLICATION: AMDER: APPLICATION: AMDER: APPLICATION: AMDER: A
                                                                                                                                                                                            FILING DATE:
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/209,204
FILING DATE: 08-MAR-1994
FRILING DATE: 08-MAR-1994
FRILING DATE: 08-MAR-1994
APPLICATION NUMBER: 08/059,022
FILING DATE: 06-MAY-93
ATTORNEY/AGENT INFORMATION:
NAME: CLARK, PAUL T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 04585/028001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
                                                                                                 SOFTWARE: Wordperfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/461,097
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-461-097A-40; Sequence 40, Application US/08461097A; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Bieker-Brady, Kristina
REGISTRATION NUMBER: 39,109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
                                                                  OPERATING SYSTEM: PC-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
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   MEDIUM TYPE:
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ZIP: 02110
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Query Match 100.0%; Score 69; DB 8; Length 39; Best Local Similarity 10.8%; Pred. No. 3.4e+03; Matches 4; Conservative 33; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:

Sequence 40, Application US/08467459

GENERAL INFORMATION:

APPLICANT: MCBURNEY, ROBERT N.

TITLE OF INVENTION: THEIR PREPARATION AND USE
NUMBER OF SEQUENCES: 184

COMESSONDECE ADDRESS:
ADDRESSE: Fish & Richardson
STREET: 225 Franklin Street
CITY: Beston
STRATE: Massachusetts
ZIP: 0211-2804
COMPUTER READBLE FORM:
MEDIUM TYBE: Diskette, 5.25 inch, 360 kb storage
COMPUTER READBLE FORM:
MEDIUM TYBE: Diskette, 5.25 inch, 360 kb storage
COMPUTER READBLE FORM:
MEDIUATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/08/467,459
FILING DATE: 29-JAN-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/984,085
FILING DATE: 07/984,085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3 CGCCKCCRTTCACRCAGAAGGTCTTCTCCTTCTCAGC 39
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INFORMATION FOR SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                      RESULT 15
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Search completed: May 27, 2005, 00:45:15 Job time : 496 secs

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NAME/KEY: misc_feature
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Sequence 26469, A
Sequence 26469, A
Sequence 53958, A
Sequence 26472, A
Sequence 2113, Ap
Sequence 9113, Ap
Sequence 26460, A
Sequence 26460, A
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Sequence 5372, Ap
Sequence 5372, Ap
Sequence 53959, A
Sequence 57805, A
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Sequence 192934,
Sequence 105601,
Sequence 29, App
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                                                                                  May 27, 2005, 00:32:31; Search time 52 Seconds (without alignments) 81.197 Million cell updates/sec
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1: /cgn2 6/ptodata/1/paa/PCT NEW COMB.pep:*

2: /cgn2 6/ptodata/1/paa/USO6 NEW COMB.pep:*

3: /cgn2 6/ptodata/1/paa/USO7 NEW COMB.pep:*

4: /cgn2 6/ptodata/1/paa/USO8 NEW COMB.pep:*

5: /cgn2 6/ptodata/1/paa/USO8 NEW COMB.pep:*

6: /cgn2 6/ptodata/1/paa/USO8 NEW COMB.pep:*

7: /cgn2 6/ptodata/1/paa/USO1 NEW COMB.pep:*

8: /cgn2 6/ptodata/1/paa/USO1 NEW COMB.pep:*
                                                                                                                                                                                                                                                                                     537409
                                                                                                                                                                             GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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US-11-097-143-26469

US-10-526-324-658

US-10-526-324-658

US-10-450-763-53958

US-11-097-143-26472

US-11-079-463-9113

US-11-079-045-9113

US-11-097-143-26460

US-11-097-143-26450

US-11-450-763-57859

US-10-450-763-57805

US-10-450-763-57805

US-10-450-763-57805

US-10-655-875-1123185

US-60-655-875-1123185
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US-10-703-032-105601
US-10-530-253-29
                                                                                                                                                                                                                                                                                    Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                      537409 seqs, 114114291 residues
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Maximum Match 100%
Listing first 45 summaries
                                                         - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 200000000
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Match Length DB
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Perfect score:
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No.
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ALIGNMENTS

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APPLICANT: Imanaka, Takayuki
APPLICANT: Imanaka, Takayuki
APPLICANT: Imanaka, Haruyuki
TITLE OF INVENTION METHOD OF TARGETED GENE DISRUPTION, GENOME OF
TITLE OF INVENTION: HYPERTHERMOSTABLE BACTERIUM AND GENOME CHIP USING
TITLE OF INVENTION: HYPERTHERMOSTABLE BACTERIUM AND GENOME CHIP USING
TITLE OF INVENTION: HYPERTHERMOSTABLE BACTERIUM AND GENOME CHIP USING
TITLE OF INVENTION: HE SAME
FILE REPRENDENCE: 40051.4011USPC
CURRENT APPLICATION NUMBER: UST/1182003/003597
PRIOR APPLICATION NUMBER: PT/1182003/003597
PRIOR PLILING DATE: 2003-08-29
PRIOR FILING DATE: 2003-08-29
NUMBER OF SEQ ID NOS: 2167
SOFTWARE: PATENTIN VERSION 3.1
SEQ ID NO 813
LENGTH: S3
LENGTH: S3
CORTANDENT S1
CORGANISM: Thermococcus kodakaraensis KOD1
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US-10-526-324-813
; Sequence 813, Application US/10526324
; GENERAL INFORMATION:
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NAME/KEY: misc_feature
LOCATION: (786907)..(786907)
OTHER INFORMATION: n is a or c or
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LOCATION: (1128488)..(1128488)
OTHER INFORMATION: n is a or c or
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LOCATION: (839139)..(839139)
OTHER INFORMATION: n is a or c or
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NAMES/KEY: misc_feature
LOCATION: (786890)
OTHER INFORMATION: n is a or
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LOCATION: (414542)..(414542)
OTHER INFORMATION: n is a or
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NAME/KEY: misc_feature
LOCATION: (786944)..(786946)
OTHER INFORMATION: n is a or
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APPLICANT: et al.

APPLICANT: et al.

TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID

TITLE OF INVENTION: ARRAYS, FOR DETECTING EXPRESSION OF 10,000 OR MORE

TITLE OF INVENTION: DROSOPHILA GENES.

FILE REFERENCE: CLO00728

CURRENT FILING DATE: 1999-10-05

PRIOR FILING DATE: 1999-10-19

PRIOR PLICATION NUMBER: 60/150,191

PRIOR PLILNG DATE: 1999-10-19

PRIOR PLILNG DATE: 1999-10-19

PRIOR APPLICATION NUMBER: 60/160,191

PRIOR PLILNG DATE: 1999-10-12

PRIOR PLILNG DATE: 1999-11-12

PRIOR PLILNG DATE: 1999-11-28

PRIOR PLILNG DATE: 2000-01-22

PRIOR PLILNG DATE: 2000-02-24

PRIOR PLILNG DATE: 2000-03-23

NUMBER: 05 SEQ ID NOS: 43008

SSOCIEMARE: FRASTSEQ for Windows Version 4.0

FROMTH. 64
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APPLICANT: Atomi, Haruyuki
TITLE OF INVENTION: METHOD OF TARGETED GENE DISRUPTION, GENOME OF
TITLE OF INVENTION: HYPBRITHERMOSTABLE BACTERIUM AND GENOME CHIP USING
TITLE OF INVENTION: THE SAME
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10.8%; Pred. No. 1.3e+02;
iive 33; Mismatches 0;
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CURRENT APPLICATION NUMBER: US/10/526,324
CURRENT FILING DATE: 2005-02-28
PRIOR APPLICATION NUMBER: PCT/IB2003/003597
PRIOR APPLICATION NUMBER: PCT/IB2003/003597
PRIOR APPLICATION NUMBER: J
                                                                                                                                                                                                                                                                                                                                                ; Sequence 26490, Application US/11097143; GENERAL INFORMATION:
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                                                                           Query Match
Best Local Similarity 10.8*
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Venter, J. Craig APPLICANT: et al.
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Best Local Similarity
Matches 4; Conserv
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; TYPE: PRT
; ORGANISM: DROSOPHILA
US-11-097-143-26469
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EXPRESSION OF 10,000 OR MORE
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ilarity 10.8%; Pred. No. 1.2e+02;
Conservative 33; Mismatches 0
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GENERAL INFORMATION:
APPLICANT: Verter, J. Craig
APPLICANT: et al.
TITLE OF INVENTION: DETECTION KIT, SUCH AS NUC:
TITLE OF INVENTION: DETECTION KIT, SUCH AS NUC:
TITLE OF INVENTION: ARRAYS, FOR DETECTING EXPITION EDGES.
TITLE OF INVENTION: DETECTION CONTROL
TITLE OF INVENTION: BARBAYS, FOR DETECTING EXPITION FOR EDGES.
FILE REFERENCE: CLOOO728
CURRENT APPLICATION NUMBER: 60/157,832
PRIOR PILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: 60/161,932
PRIOR PILING DATE: 1999-10-28
PRIOR PILING DATE: 1999-11-12
PRIOR PILING DATE: 1999-11-2
PRIOR APPLICATION NUMBER: 60/161,932
PRIOR APPLICATION NUMBER: 60/161,933
PRIOR PILING DATE: 1999-11-2
PRIOR APPLICATION NUMBER: 60/175,693
PRIOR PILING DATE: 2000-01-12
PRIOR PILING DATE: 2000-01-12
PRIOR PILING DATE: 2000-03-23
NUMBER: 60/191,637
PRIOR PILING DATE: 2000-03-23
NUMBER: FSEAUSEQ fOR WINDOWS VERSION 4.0
SEQ ID NO 26469
LENGTH: 55
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LOCATION: (1561477)..(1561477)
OTHER_INFORMATION: n is a or c or g or
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LOCATION: (1128517)..(1128518)
OTHER INFORMATION: n is a or c or g or
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                               OTHER INFORMATION: n is a or c or g or
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                                                                      NAME/KEY: misc_feature
LOCATION: (1128505)..(1128506)
OTHER INFORMATION: n is a or c or g
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LOCATION: (1561400)..(1561400)
OTHER INFORMATION: n is a or c
FEATURE:
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LOCATION: (1561545)..(1561545)
OTHER INFORMATION: n is a or c
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LOCATION: (1128539)..(1128540)
OTHER INFORMATION: n is a or c
       LOCATION: (1128499) .. (1128499)
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Best Local Similarity
Matches 4; Conserv
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LOCATION: (10)..(56)
OTHER INFORMATION: Vertebrate metallothioneins proteins domain identified by
OTHER INFORMATION: eMATRIX, accession number BL00203, p-value=9.603e-21, raw score
OTHER INFORMATION: 13.94
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TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID

TITLE OF INVENTION: DROSOPHILA GENES.

TITLE OF INVENTION: DROSOPHILA GENES.

TITLE OF INVENTION: DROSOPHILA GENES.

CURRENT APPLICATION NUMBER: US/11/097,143

CURRENT APPLICATION NUMBER: 60/157,832

PRIOR PILING DATE: 1999-10-05

PRIOR PILING DATE: 1999-10-19

PRIOR PILING DATE: 1999-10-19

PRIOR PILING DATE: 1999-11-12

PRIOR PILING DATE: 1999-11-2

PRIOR PILING DATE: 2000-01-12

PRIOR PILING DATE: 2000-01-12

PRIOR PILING DATE: 2000-01-2

PRIOR PILING DATE: 2000-01-2

PRIOR PILING DATE: 2000-01-2

PRIOR PILING DATE: 2000-01-3

PRIOR PILING DATE: 2000-
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                                                                                                                                           APPLICANT: Hyeeq, Inc.
TITLE OF INVENTION: NOVEL NUCLEIC ACIDS AND POLYPEPTIDES
FILE REFERENCE: 790CIP3/US
CURRENT APPLICATION NUMBER: US/10/450,763
CURRENT APPLICATION NUMBER: US/10/631
PRIOR ALPLICATION NUMBER: PCT/US01/08631
PRIOR FILING DATE: 2001-03-30
PRIOR FILING DATE: 2000-03-31
PRIOR PILING DATE: 2000-03-31
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR PILING DATE: 2000-03-31
PRIOR PILING DATE: 2000-03-31
PRIOR PILING DATE: 2000-03-31
NUMBER OF SEQ ID NOS: 60736
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SOFTWARE: FastSEQ for Windows Version 4.0
                                RESULT 5
US-10-450-763-53958
US-10-450-763-53958
Sequence 53958, Application US/10450763
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Homo sapiens
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ORGANISM: DROSOPHILA
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LOCATION: (10)...
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LENGTH: 63
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                                                                                           SEQ ID NO 658
LENGTH: 59
TYPE: PRT
ORGANISM: Thermococcus kodakaraensis KOD1
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LOCATION: (1561477)...(1561477)
OTHER INFORMATION: n is a or c or g or t.
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LOCATION: (1561545)..(1561545)
OTHER INFORMATION: n is a or c or g or t.
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LOCATION: (414542)..(414542)
OTHER INFORMATION: n is a or c or g or t.
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LOCATION: (1128505)..(1128506)
OTHER INFORMATION: n is a or c or g or t.
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LOCATION: (839139)..(839139)
OTHER INFORMATION: n is a or c or g or
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LOCATION: (786944)..(786946)
OTHER INFORMATION: n is a or c or g or
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LOCATION: (786907)..(786907)
OTHER INFORMATION: n is a or c or g
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LOCATION: (1128488)..(1128488)
OTHER INFORMATION: n is a or c or g
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LOCATION: (1128517)..(1128518)
OTHER INFORMATION: n is a or c or g
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LOCATION: (1128539)..(1128540)
OTHER INFORMATION: n is a or c or g
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LOCATION: (1561400)..(1561400)
OTHER INFORMATION: n is a or c or g
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LOCATION: (786890)..(786890)
OTHER INFORMATION: n is a or c or g
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LOCATION: (1128499)..(1128499)
OTHER INFORMATION: n is a or c
PRIOR FILING DATE: 2002-08-30
NUMBER OF SEQ ID NOS: 2167
SOFTWARE: PatentIn version 3.1
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Best Local Similarity
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Matches
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTEROIDES FR
TITLE OF INVENTION: NUCLEIC ACID AND THERAPEUTICS
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REPERBACE: PATHOD-03DIV1
CURRENT APPLICATION NUMBER: US/11/079,045
PRIOR FILING DATE: 1099-04-09
PRIOR FILING DATE: 1099-04-09
PRIOR FILING DATE: 2000-04-04
NUMBER OF SEQ ID NOS: 10444
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TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID
TITLE OF INVENTION: ARRAYS, FOR DETECTING EXPRESSION OF 10,000 OR MORE
TITLE OF INVENTION: DECOSOPHILA GENES.
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23 CIVCDFIYDPEIGDPEGGIEPGTOFEDIPDDWVCPLC 59
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Best Local Similarity 10.8%; Pred. No. 1.8e+02;
Matches 4; Conservative 33; Mismatches 0.
                                                                                                  100.0%; Score 69; DB 7; 10.8%; Pred. No. 1.8e+02;
                                                                                                                                            33, Mismatches
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US-11-097-143-26460
; Sequence 26460, Application US/11097143
; GENERAL INFORMATION:
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GENERAL INFORMATION:
                                                                                   Query Match
Best Local Similarity 10.0%
The At Conservative
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US-11-079-045-9113
                 ; TYPE: PRT
; ORGANISM: B.fragilis
US-11-079-463-9113
                                                                                                                                                                                                                                                                                                                   US-11-079-045-9113
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LENGTH: 71
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FILE REFERENCE: 38-21(53.596)
CURRENT APPLICATION NUMBER: US/60/669,241
CURRENT FILING DATE: 2005-04-07
PRIOR APPLICATION NUMBER: 6056.0842
PRIOR APPLICATION NUMBER: 6056.032
PRIOR APPLICATION NUMBER: 6056.032
PRIOR FILING DATE: 2004-04-27
PRIOR PRIOR DATE: 2004-04-27
PRIOR APPLICATION NUMBER: 6057.9062
PRIOR APPLICATION NUMBER: 6060.3421
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Sequence 9113, Application US/11079463
GENERAL INFORMATION:
APPLICANT: Gary L. Breton
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTEROIDES FRA
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: PATHO0-03DIV2
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US-60-669-241-25327
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                                       Query Match 100.0%; Score 69; DB 7; Length 63; Best Local Similarity 10.8%; Pred. No. 1.5e+02; Matches 4; Conservative 33; Mismatches 0; Indels
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                                                                                                                                                     26 CGGCGPCCGGCGPCCGPCCGGTSSFCGCGPC 62
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CURRENT FILING DATE: 2005-03-14
PRIOR APPLICATION NUMBER: US 60/128,705
PRIOR PILING DATE: 1999-04-09
PRIOR APPLICATION NUMBER: US 09/540,209
PRIOR APPLICATION NUMBER: US 09/540,209
PRIOR FILING DATE: 2000-04-04
NUMBER OF SEQ ID NOS: 10444
                                                                                                                                                                                                                                                                         Sequence 25327, Application US/60669241 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                       Baum, James A
Gilberteson, Larry A
Kovalic, David K
LaRosa, Thomas J
Lu, Maolong
Munnikwa, Tichifa R. I.
Roberts, James K
                                                                                                                                                                                                                                                                                                               APPLICANT: Monsanto Technology LLC
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US-11-097-143-26472
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LENGTH: 66
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APPLICANT:
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US-60-669-175-32704
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APPLICANT: Conner, Timothy W.
APPLICANT: Cao, Yongwei
APPLICANT: Masucci, James D.
APPLICANT: Zhou, Yihua
TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With
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APPLICANT: et al.
TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID
TITLE OF INVENTION: ARRAYS, FOR DETECTING EXPRESSION OF 10,000 OR MORE
TITLE OF INVENTION: DROSOPHILA GENES.
TITLE OF INVENTION: USOSOPHILA GENES.
CURRENT APPLICATION NUMBER: US/11/097,143
CURRENT APPLICATION NUMBER: 2005-04-04
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                                                                                                                                              100.0%; Score 69; DB 7; Length 72; larity 10.8%; Pred. No. 1.8e+02; Conservative 33; Mismatches 0; Indels
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Pred. No. 1.9e+02;
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PRIOR APPLICATION NUMBER: 60/157,832

PRIOR FILING DATE: 1999-10-05

PRIOR FILING DATE: 1999-10-19

PRIOR PILING DATE: 1999-10-19

PRIOR PILING DATE: 1999-10-19

PRIOR PLING DATE: 1999-10-28

PRIOR PLING DATE: 1999-11-12

PRIOR APPLICATION NUMBER: 60/164,769

PRIOR FILING DATE: 1999-11-12

PRIOR PLING DATE: 1999-11-28

PRIOR FILING DATE: 1999-11-28

PRIOR FILING DATE: 1999-11-28

PRIOR FILING DATE: 1900-01-28

PRIOR PRIOR DATE: 2000-01-28

PRIOR PRIOR DATE: 2000-01-12

PRIOR APPLICATION NUMBER: 60/191,637

PRIOR PRIING DATE: 2000-02-24

PRIOR PRIING DATE: 2000-03-23

NUMBER: FASTERD FOR WINDGER: 60/191,637

PRIOR FILING DATE: 2000-03-23

NUMBER: FASTERD FOR WINDGER: 60/191,637

PRIOR FILING DATE: 2000-03-23

NUMBER: FASTERD FOR WINDGER: 60/191,637
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SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 26460
LENGTH: 72
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                                                                                                                                                                                                                                                                                                                                                                             Sequence 26457, Application US/11097143 GENERAL INFORMATION:
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10.8%; F
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APPLICANT: Andersen, Scott E.
APPLICANT: Byrum, Joseph R.
APPLICANT: Conner, Timothy W.
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                                                                                                                                            Query Match
Best Local Similarity
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US-10-703-032-107111
                                                                                                     US-11-097-143-26460
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LENGTH: 74
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APPLICANT: Cao, Yongwei
APPLICANT: Cao, Yongwei
APPLICANT: Masucci, James D.
APPLICANT: Macucci, James D.
APPLICANT: Abou, Yihua
TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(53374)8
CURRENT APPLICATION NUMBER: US/10/703,032
PRIOR APPLICATION NUMBER: 10/020,338
PRIOR PLING DATE: 2001-12-12
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100.0%; Score 69; DB 6, Best Local Similarity 10.8%; Pred. No. 2e+02; Matches 4; Conservative 33; Mismatches
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OTHER INFORMATION: unsure at all Xaa locations
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100.0%; Score 69; DB 6
Best Local Similarity 13.5%; Pred. No. 2e+02;
Matches 5; Conservative 32; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: Clone ID: PAT_TA_86449.pep
US-10-703-032-192031
FILE REPERENCE: 38-21(53374)B
; CURRENT APPLICATION NUMBER: US/10/703,032
; CURRENT FILING DATE: 2003-11-06
; PRIOR PALLICATION NUMBER: 10/020,338
; PRIOR PLIING DATE: 2001-12-12
; NUMBER OF SEQ ID NOS: 211164
; SEQ ID NO 107111
; LENGTH: 76
                                                                                                                                                                                                                                                                                                       ; OTHER INFORMATION: Clone ID: PAT_TA_1529.pep
US-10-703-032-107111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 192031, Application US/10703032 GENERAL INFORMATION:
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Gilbertson, Larry A
Kovalic, David K
LaRosa, Thomas J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Kovalic, David K.
APPLICANT: Andersen, Scott E.
APPLICANT: Byrum, Joseph R.
APPLICANT: Conner, Timothy W.
                                                                                                                                                                                                                                        TYPE: PRT ORGANISM: Triticum aestivum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Triticum aestivum
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SEQ ID NO 192031
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APPLICANT:
APPLICANT:
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APPLICANT: Zhang, Bei
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR CONTROL OF INSECT INFESTATION IN PLA
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR CONTROL OF INSECT INFESTATION IN PLA
TILE REPERENCE: 38-21 (53597)
CURRENT APPLICATION NUMBER: 0505-04-07
PRIOR APPLICATION NUMBER: 6056042
PRIOR FILING DATE: 2004-04-09
PRIOR FILING DATE: 2004-04-07
PRIOR APPLICATION NUMBER: 6056322
PRIOR APPLICATION NUMBER: 60603421
PRIOR PILING DATE: 2004-06-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TYPE: PRT
; ORGANISM: Diabrotica virgifera
; OTHER INFORMATION: Coding regions on vcDNA: vcDNA=SeqID_12401; Strand=-; Position=59
US-60-669-175-32704
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CURRENT PAPLICATION NUMBER: US/10/467,657
CURRENT FILING DATE: 2003-08-11
PRIOR APPLICATION NUMBER: GB-0103424.8
PRIOR FILING DATE: 2001-02-12
NUMBER OF SEQ ID NOS: 9218
SOFTWARE: SEQ ID NOS: 9218
SOFTWARE: SEQ ID NOS: 9218
SOFTWARE: SEQ ID NOS: 9218
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Job time : 53 secs
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; Sequence 5372, Application US/10467657
; GENERAL INFORMATION:
Lu, Maolong
Munyikwa, Tichifa R. I.
Roberts, James K
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; ORGANISM: Neisseria gonorrhoeae
US-10-467-657-5372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: CHIRON SpA
APPLICANT: FONTANA Maria Rita
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: PIZZA Mariagrazia
APPLICANT: MASIGNANI Vega
APPLICANT: MONACI Elisabetta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 40774
SEQ ID NO 32704
LENGTH: 80
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Best Local Similarity
Matches 4; Conserv
                                                                           APPLICANT:
APPLICANT:
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